

Title

**Summary of the Literature Review for GHB614 x LLCotton25 x MON 15985 Cotton
July 1, 2022 – June 30, 2023**



Final ReportData or guideline requirement

Explanatory note on literature searching
conducted in the context of GMO applications for (renewed) market authorization
and annual post-market environmental monitoring reports on GMOs authorised in the EU market.
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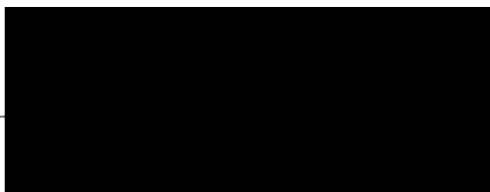
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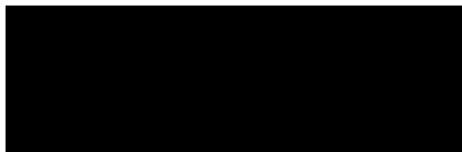
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SUMMARY

BASF has used conventional breeding techniques to develop the stacked trait cotton product GHB614 x LLCotton25 x MON 15985 (GLB2 cotton) which confers resistance to lepidopteran insects and tolerance to glyphosate and glufosinate-ammonium herbicides. The OECD identifier is BCS-GHØØ2-5 x ACS-GHØØ1-3 x MON-15985-7.

A scoping review was performed for the GHB614 x LLCotton25 x MON 15985 cotton and its newly expressed proteins, 2mEPSPS, PAT/*bar*, Cry1Ac, and Cry2Ab2. The objective of this scoping review was to determine if there were studies about the molecular characterization of GHB614 x LLCotton25 x MON 15985 cotton, its effect on food and feed safety or environmental safety, that might require in-depth examination. A set of broad literature searches was performed using several bibliographic databases covering scientific literature from July 1, 2022 to June 30, 2023. Additional sources of information, such as web pages of food safety, agriculture, and biotechnology-related authorities were searched for the same time window, along with the bibliographies of relevant reviews. The references identified were evaluated for potential relevance to the scoping review questions according to pre-defined criteria.

These literature searches identified a total of 49 unique publications, which were subject to rapid assessment to exclude obviously irrelevant publications. Five publications were progressed for detailed assessment and determined to be not relevant after detailed review.

No new publications were found that contained new data on the molecular characterization of the GHB614 x LLCotton25 x MON 15985 cotton and its newly expressed protein, 2mEPSPS, PAT/*bar*, Cry1Ac, and Cry2Ab2. Similarly, no new publications were found that suggested any potential adverse effects of GHB614 x LLCotton25 x MON 15985 cotton on human health, animal health, or the environment. No issues or topics were identified that would trigger or warrant more specific question formulation or indicate that a systematic review would be of value.

1. INTRODUCTION

BASF has used conventional breeding techniques to develop the stacked trait cotton product GHB614 x LLCotton25 x MON 15985 (GLB2 cotton) which confers resistance to lepidopteran insects and tolerance to glyphosate and glufosinate-ammonium herbicides. The OECD identifier is BCS-GHØØ2-5 x ACS-GHØØ1-3 x MON-15985-7.

The objective of the literature searches described here was to determine if there were publications published between July 1, 2022 and June 30, 2023 that mention the molecular characterization of the GHB614 x LLCotton25 x MON 15985 cotton, and/or any adverse effect of GHB614 x LLCotton25 x MON 15985 cotton in food, feed or the environment. In that context, a broad and inclusive literature search was performed, and the articles retrieved were reviewed in a comprehensive and transparent manner. This was intended as a scoping review. The literature review was performed as recommended in the European Food Safety Authority (EFSA) explanatory note on literature searching conducted in the context of Genetically Modified Organisms (GMO) applications and post-market environmental monitoring activities (2019, (1)).

The literature searches were performed for the GHB614 x LLCotton25 x MON 15985 cotton and its newly expressed proteins, 2mEPSPS, PAT/*bar*, Cry1Ac, and Cry2Ab2. The search terms also included relevant synonyms, the trade name and intended trait. When needed, plant species and general GMO terms were used to limit the search results (described in [Section 3.3](#)).

2. OVERALL METHODS

2.1. Objective of the scoping review

The objective of the scoping review was to survey the evidence base for the GHB614 x LLCotton25 x MON 15985 cotton and its newly expressed proteins, 2mEPSPS, PAT/*bar*, Cry1Ac, and Cry2Ab2, in order to identify any specific issues related to food or feed safety, molecular characterization or environmental safety that might require in-depth examination.

2.2. Review questions

Review questions were formulated to conform to PE(I)CO structure (Population, Exposure (Intervention), Comparators, Outcome) if possible, and to address data requirements. They were modeled after the review question examples provided in the EFSA 2019 explanatory note (1).

Question 1: Were any studies published during the reporting period that describe adverse effects on human or animal health or the environment of the GHB614 x LLCotton25 x MON 15985 cotton and its newly expressed proteins 2mEPSPS, PAT/*bar*, Cry1Ac, and Cry2Ab2?

Key elements:

Population: Human health; animal health; environmental safety

Exposure: GHB614 x LLCotton25 x MON 15985 cotton, derived food/feed products, newly expressed proteins in GHB614 x LLCotton25 x MON 15985 cotton

Comparators: When applicable, comparable populations or subjects exposed to appropriate controls (e.g., vehicle only, innocuous control protein, non-GM comparator) or conventional counterpart used for comparative analysis of plant material

Outcome: Adverse effects

Question 2: Were any studies published during the reporting period that focus on molecular characterization of the GHB614 x LLCotton25 x MON 15985 cotton and its newly expressed proteins 2mEPSPS, PAT/*bar*, Cry1Ac, and Cry2Ab2 in cotton?

Key elements:

Population: GHB614 x LLCotton25 x MON 15985 cotton and newly expressed proteins in GHB614 x LLCotton25 x MON 15985 cotton

Outcome: Molecular characterization (which would indicate the information/data requirement for molecular characteristics)

2.3. Criteria for relevance

Criteria for establishing the relevance of retrieved publications were defined prior to conduct of the search. These criteria were modeled after those given in the EFSA 2019 explanatory note (1) and are described in Table 1.

Table 1: Eligibility/inclusion criteria to establish the relevance of retrieved publications

Concepts	Criteria	Comment
Key elements of review questions with PECO structure		
Population	The publication addresses human and animal health, and/or the environment (including biodiversity, ecosystem services, service providing units, and endangered species) as general protection goals	From the publications that address the GMO under consideration, those that address protection goals relevant to the risk assessment of the GMO are eligible
Exposure (Intervention)	The publication addresses the GMO, derived food/feed products, and/or the intended trait(s) (e.g., newly expressed proteins(s)) that are identical or like those under regulatory review	This enables the selection of publications that address the GMO, derived food/feed products, and/or the intended trait(s) under consideration
Comparator	If the publication reports a comparative study that uses plant material as test material, eligible publications must report a non-GM variety as comparator	In those cases where the publication addresses the GMO under consideration, reports a comparative analysis study and uses plant material as test material, eligible publications also need to include an appropriate non-GM line as comparator
Outcome	The publication addresses effects/impacts on human and animal health, and/or the environment	Publications that address the GMO under consideration also need to address effects/impacts on entities of concern, and potential determinants of exposure that place these entities at risk, in order to be relevant to the risk assessment of the GMO

Concepts	Criteria	Comment
Additional concepts		
Information/data requirements	The publication reports information pertaining to one or more information/data requirement(s) outlined in Appendix A for the GMO and derived food/feed products under consideration, including the intended trait(s)	Publications that potentially contribute to the knowledge informing the risk assessment of the GMO under consideration, and thus the risk hypotheses addressed, taking account of both hazard and exposure, can be considered relevant according to this eligibility/inclusion criterion. Publications addressing other issues such as benefits, socio-economics, ethics, crop protection, detection methods, efficacy, public perception and risk communication can be excluded, as they are not necessarily relevant to the risk assessment of GMOs
Plant species	The publication addresses the same plant species as the GMO under consideration	This eligibility/inclusion criterion permits the exclusion of publications on GMOs that contain the same intended trait(s) as the GMO under consideration, but which are introduced in another plant species
Scope of GMO application	The publication addresses pathways and levels of exposure to the GMO, derived food/feed products, and the intended trait(s) that are relevant for the intended uses of the GMO and derived food/feed products under regulatory review	From the publications that address the GMO under consideration, those that consider pathways and levels of exposure relevant to the scope of the GMO application (<i>i.e.</i> , import and processing for food/feed uses, cultivation) are eligible
Target pests/organisms	The publication addresses target pests/organisms that are established in the EU	This permits the exclusion of publications that address interactions between the GMO and target pests/organisms that do not occur in the EU

Concepts	Criteria	Comment
Stacked events obtained by conventional crosses/ subcombinations	The publication addresses the higher stacked event and/or a subcombination or subcombinations of the single events of the higher stacked event, independently of its/their origin	This permits the selection of publications on the higher stacked event and/or subcombinations of the single events of the higher stacked event that are in the scope of the GMO application(s), independently of their origin. This permits the exclusion of publications on the single events of the higher stacked event, because the risk assessment of GMO applications for stacked events covers only the products in the scope of the GMO application – <i>i.e.</i> , the higher stacked event and subcombinations of the singles involved, independently of their origin
Molecular stacks	The publication addresses: the molecular stack; all newly expressed proteins in the molecular stack; and/or one or several of the newly expressed proteins in the molecular stack that has/have not been previously risk assessed by EFSA and/or its GMO Panel and for which no safe use has been determined yet by EFSA and/or its GMO Panel	This permits the exclusion of publications that address one or several (not all) of the newly expressed proteins in the molecular stack that has/have been previously risk assessed by EFSA and/or its GMO Panel and for which the safe use has been determined by EFSA and/or its GMO Panel
Previously risk assessed publications	The publication has not been previously risk assessed by EFSA and/or its GMO Panel and is not cited/referenced in an EFSA/GMO Panel output	This permits the exclusion of publications that have been previously risk assessed by EFSA and/or its GMO Panel and cited/referenced in an EFSA/GMO Panel output
Access	Full-text document is accessible	If potentially relevant full-text documents cannot be obtained, they should be listed in a table with a description of the (unsuccessful) methods that have been used to try to obtain a copy

Concepts	Criteria	Comment
Reporting format	The publication presents original/primary data, or it is a risk assessment from a relevant key organisation (such as regulatory agencies and risk assessment bodies involved in the risk assessment of GMOs)	This permits the exclusion of publications that do not present original/primary data (e.g., editorials, position papers), and the inclusion of relevant risk assessments performed and reported by relevant key organisations. Reviews should only be included if they present data that are not available from a primary research study
Reporting format	A study in a publication should only be presented once, but if it is presented in more than one publication, all publications should be listed and grouped	Duplicate publications should be excluded at the screening stage. Only one copy of a study is required even if it is reported in different publications, and identified in more than one database

Table adapted from EFSA, 2019: Explanatory note on literature searching conducted in the context of GMO applications for (renewed) market authorisation and annual post-market environmental monitoring reports on GMOs authorised in the EU market (1).

2.4. Reference publication

Two publication related to GHB614 x LLCotton25 x MON 15985 cotton stack were previously identified and used to test and validate the search strategy:

- Naegeli, H.; Birch, A. N.; Casacuberta, J.; Schrijver, A. de; Gralak, M. A.; Guerche, P.; Jones, H.; Manachini, B.; Messean, A.; Nielsen, E. E.; Nogue, F.; Robaglia, C.; Rostoks, N.; Sweet, J.; Tebbe, C.; Visioli, F.; Wal, J. M.; Broll, H.; Gennaro, A.; Neri, F. M.; Paraskevopoulos, K.; de Schrijver, A. (2018). Assessment of genetically modified cotton GHB614 .times. LLCotton25 .times. MON 15985 for food and feed uses, under Regulation (EC) No 1829/2003 (application EFSA-GMO-NL-2011-94). EFSA Journal 16(4): e05213 p.
- Assif, M.; Siddiqui, H.A.; Naqvi, R. Z.; Amin, I.; Asad, S.; Mukhtar, Z.; Bashir, A.; Mansoor. S. (2021). Development of event-specific detection method for identification of insect resistant NIBGE-1601 cotton harboring double gene Cry1Ac-Cry2Ab construct. Scientific Reports 11, article 3479.

The first article is directly relevant for the GHB614 x LLCotton25 x MON 15985 cotton stack, while the second article mentions two of the newly expressed proteins (Cry1Ac and Cry2Ab), the crop (cotton) and one of the intended traits (insect resistance). Since these references were published before the current search period, the search profile was tested without applying the time limit used in the final search profile (UP>=20220701 and UP<=20230630).

3. SEARCH METHODS AND OUTCOMES

The search strategies used here followed the 2019 EFSA explanatory note on literature searching conducted in the context of GMO applications and post-market environmental monitoring activities (1). The search strategies were designed to be broad and sensitive enough to capture any relevant publications, if available.

An information specialist with background in plant biotechnology selected the databases, identified relevant search terms, developed search profiles, designed search strategies, and conducted the searches.

3.1. Time window and date of the literature search

The database searches were performed on July 14, 2023. Only documents updated between July 1, 2022 and June 30, 2023, were considered in the search. The dates of most recent database updates are provided in [Table 4](#).

3.2. Databases used in the literature search

All searches were performed in the host STN (Scientific and Technical Information Network), an online database service operated jointly by CAS and FIZ Karlsruhe. STN provides access to a broad range of databases from the most renowned database producers worldwide.

The searches described here were performed in five databases: three multidisciplinary/large databases (Biosis, Medline and CA-Plus) and two subject-specific databases focused on agriculture-related topics (Agricola and CABA).

See [Appendix 1](#) for detailed database descriptions.

3.3. Search strategy

The search profiles were designed to cover event name, newly expressed proteins and intended traits of the stack. Since the 'intended traits' profile identified too many references when used alone, it was combined with a 'general GMO' profile and a 'plant species' profile. The reference publication ([Section 2.4](#)) was identified by the search profiles confirming the validity of the applied search strategy. See Table 2 for a detailed search profile.

Table 2: Search profile for database search

Set	Search string	Concepts
1	GHB614 OR GHB(W)614 OR BCS-GH002-5 OR BCSGH002-5 OR BCS(W)GH002(W)5 OR BCSGH002(W)5 OR BCS-GH002-5 OR BCS(W)GH002(W)5 OR BCSGH002(W)5 or GHB614x OR GHB(W)614x OR BCS-GH002-5x OR BCSGH002-5x OR BCS(W)GH002(W)5x OR BCSGH002(W)5x OR BCS-GH002-5x OR BCS(W)GH002(W)5x OR BCSGH002(W)5x	Event name GHB614
2	LLcotton25 or LLcotton(w)25 or LL(w)cotton25 or LL(w)cotton(w)25 or ACS-GH001-3 or ACS(w)GH001(w)3 or ACSGH001(w)3 or ACS-GH001-3 or ACS(w)GH001(w)3 or ACSGH001(w)3 or LLcotton25x or LLcotton(w)25x or LL(w)cotton25x or LL(w)cotton(w)25x or ACS-GH001-3x or ACS(w)GH001(w)3x or ACSGH001(w)3x or ACS-GH001-3x or ACS(w)GH001(w)3x or ACSGH001(w)3x or xLLcotton25	Event name LLCotton25

Set	Search string	Concepts
	or xLLcotton(w)25 or xLL(w)cotton25 or xLL(w)cotton(w)25 or xACS-GH001-3 or xACS(w)GH001(w)3 or xACSGH001(w)3 or xACS-GH001-3 or xACS(w)GH001(w)3 or xACSGH001(w)3 or xLLcotton25x or xLLcotton(w)25x or xLL(w)cotton25x or xLL(w)cotton(w)25x or xACS- GH001-3x or xACS(w)GH001(w)3x or xACSGH001(w)3x or xACS-GH001-3x or xACS(w)GH001(w)3x or xACSGH001(w)3x	
3	MON(w)15985 or MON15985 or MON-15985-7 or MON(w)15985(w)7 or MON15985(w)7 or xMON(w)15985 or xMON15985 or xMON-15985-7 or xMON(w)15985(w)7 or xMON15985(w)7	Event name MON 15985
4	GHB614XLLcotton25xMON15985 or GHB(w)614XLLcotton25xMON15985 or GHB614XLL(w)cotton25xMON15985 or GHB614XLLcotton(w)25xMON15985 or GHB614XLLcotton25xMON(w)15985 or GHB(w)614XLL(w)cotton25xMON15985 or GHB(w)614XLLcotton(w)25xMON15985 or GHB(w)614XLLcotton25xMON(w)15985 or GHB(w)614XLL(w)cotton(w)25xMON15985 or GHB(w)614XLL(w)cotton25xMON(w)15986 or GHB(w)614XLL(w)cotton(w)25xMON(w)15985 or GHB614(w)time#(w)LLcotton25(w)time#(w)MON15985 or GHB614(w)time#(w)LLcotton25(w)time#(w)MON(w)15985	Event name stack
5	(1 and 2) or (1 and 3) or (2 and 3) or 4	Event name all
6	GLYTOL? OR GLYTOLTM? OR GLYTOLRTM? OR GLY(w)TOL? OR GLY(w)TOLTM? OR GLY(w)TOLRTM?	Trade name GHB614
7	libertylink or liberty(w)link or libertylinktm or liberty(w)linktm or libertylinkrtm or liberty(w)linkrtm	Trade name LLCotton25
8	bollgard? or bolgard?	Trade name MON 15985
9	(6 and 7) or (6 and 8) or (7 and 8)	Trade name all
10	(2MEPSPS or 2(w)MEPSPS or 2M(w)EPSPS or 2(w)M(w)EPSPS) or ((EPSPS OR EPSP(W)SYNTHASE OR (ENOL(W)PYRUVYLSHIKIMATE OR ENOL(W)PYRUVYL(W)SHIKIMATE OR ENOLPYRUVYLSHIKIMATE OR ENOLPYRUVYOYLSHIKIMATE or ENOYLPYRUVYOYLSHIKIMATE OR ENOLPYRUVYLSHIKIMIC) (4W) (PHOSPHATE OR PHOSPHORIC) (2W) (SYNTHASE OR SYNTHETASE) or (ENOLPYRUVYL OR ENOLPYRUYL OR ENOLPYRUVOYL) (W) (PHOSPHOSHIKIMATE OR PHOSPHOSHIKIMIC or ENOLPYRUVYLSHIKIMATEPHOSPHATE) (2W) (SYNTHASE OR SYNTHETASE) or (ENOL(W)PYRUVYOYLSHIKIMATE OR ENOLPYRUVYLSHIKIMATE OR ENOLPYRUVYLSHIKIMIC OR ENOL(W) (PYRUVYL OR PYRUVOYL) (W) SHIKIMATE) (3W) PHOSPHATE (W) (SYNTHASE OR SYNTHETASE) or (PHOSPHOSHIKIMATE (2W) CARBOXYVINYLTRANSFERASE OR PHOSPHOSHIKIMATE (2W) CARBOXYVINYL (W) TRANSFERASE OR ENOLPYRUVOYL (W) SHIKIMIC (3W) PHOSPHOSYNTHASE) (s) ((DOUBL# or DOBL#) (W) (MUTANT# OR MUTAT?) OR 2M))	Newly expressed proteins GHB614
11	((bar or pat) (2a) (gene# or protein# or enzyme#)) or ppt (2w) acetyltransferase or	Newly expressed proteins LLCotton25

Set	Search string	Concepts
	ppt(2w)acetyl(w)transferase or pt(w)n(2w)acetyltransferase or pt(w)n(2w)acetyl(w)transferase or phosphinothricin(w)n(w)acetyltransferase or phosphinothricin(2w)acetyltransferase or phosphinothricin(2w)acetyl(w)transferase or phosphinothricinacetyl(w)transferase	
12	crylaC# or cry(w)l(w)aC# or cry(w)laC# or cryl(w)aC# or cryl(w)a(w)c# or cry(w)l(w)a(w)c# or cryla(w)c# or cryIaC# or cry(w)I(w)aC# or cry(w)IaC# or cryI(w)aC# or cryI(w)a(w)c# or cry(w)I(w)a(w)c# or cryIa(w)c# or crylaC# or cry(w)l(w)aC# or cry(w)laC# or cryl(w)aC# or cryl(w)a(w)c# or cry(w)l(w)a(w)c# or cryla(w)c# or Cry2Ab2 or Cry(w)2Ab2 Cry2(w)Ab2 or Cry2A(w)b2 or Cry2Ab(w)2 or Cry2(w)Ab(w)2 or Cry(w)2(w)Ab2 or Cry(w)2(w)A(w)b2 or CryiiAb2 or Cry(w)iiAb2 Cryii(w)Ab2 or CryiiA(w)b2 or CryiiAb(w)2 or Cryii(w)Ab(w)2 or Cry(w)ii(w)Ab2 or Cry(w)ii(w)A(w)b2	Newly expressed proteins MON 15985
13	(10 and 11) or (10 and 12) or (11 and 12)	Newly expressed proteins all
14	(herbicid? or GL!PHOSATE# or GL!FOSATE# OR G360 or g(w)360 or roundup? or round(w)up?) (5a) (resist? or toleran? or protect?)	Intended traits GHB614
15	(herbicid? or bialaphos or basta or glufosinate or gluphosinate or phosphinothricin or liberty?) (5a) (resist? OR protect? OR toleran?)	Intended traits LLCotton25
16	(Insect# OR pest# OR Lepidoptera# OR Noctuidae OR Crambidae OR borer# OR cornborer# OR stalkborer# OR earworm# OR ear(w)worm# OR armyworm# OR army(w)worm# OR cutworm# OR cut(w)worm# OR Ostrinia OR O(w)nubilalis OR Sesamia OR S(w)nonagrioides or Diatraea OR D(w)grandiosella OR D(w)crambidoides OR Helicoverpa OR H(w)zea OR Spodoptera OR S(w)frugiperda OR Papaipema OR P(w)nebris OR Elasmopalpus OR E(w)lignosellus OR D(w)saccharalis OR Striacosta OR S(w)albicosta or Agrotis OR A(w)ipsilon OR S(w)cretica OR Mythimna OR M(w)unipuncta OR ECB OR MCB OR SWCB OR SCSB OR CEW OR FAW OR SCB OR WBC) (5a) (resist? OR protect? OR toleran?)	Intended traits MON 15985
17	(14 and 15) or (14 and 16) or (15 and 16)	Intended traits all
18	cotton# or gossypium or G(w)hirsutum or g(w)barbadense	Plant species
19	GMO OR GMOs OR LMO OR LMOs OR GM OR GE OR transgen? OR (genetic?(3a)(modif? OR transform? OR manipulat? OR improv? OR engineer?))	GMO general
20	17 and 18 and 19	Intended traits all AND Plant species AND GMO general
21	5 or 9 or 13 or 20	Event name all OR Trade name all OR Newly expressed proteins all species) OR (Intended

Set	Search string	Concepts
		traits all AND Plant species AND GMO general)

All searches were performed in the Basic Index (BI) field, which includes the following subject headings/field names:

- **Agricola:** title (TI), controlled term (CT), supplementary term (ST), abstract (AB), named person (NA), corporate name (CO), note (NTE), geographic term, CABA and other fields (GT)
- **Biosis:** title (TI), abstract (AB), biosystematic codes (BC), chemical name (CN), controlled term (CT), gene name (GEN), geographic term (GT), organism (ORGN) and supplementary term (ST); as well as CAS Registry Numbers (RN)
- **CA-Plus:** title (TI), supplementary term (ST), index term (IT) and abstract (AB); as well as CAS Registry Numbers
- **CABA:** title (TI), controlled term (CT), supplementary term (ST), broader term (BT), abstract (AB), organism name (ORGN) and geographic term (GT); as well as CAS Registry Numbers
- **Medline:** title (TI), chemical name (CN), gene name (GEN), controlled term (excluding MeSH numbers) (CT), supplementary term (ST), named person (NA), other source (OS), and abstract (AB), as well as CAS Registry Numbers and GenBank Numbers

Relevant controlled terms (Table 3) were not searched separately because they are included in the Basic Index and were captured by the free-text searches.

Table 3: Relevant controlled terms (CT) and index terms (IT) in each database

Database	Event	New proteins	Intended traits	Plant species	GM plants
Agricola	None	None	"HERBICIDE RESISTANCE" No terms for insect resistance	"GOSSYPIMUM BARBADENSE" "GOSSYPIMUM HIRSUTUM"	"TRANSGENIC PLANTS"
Biosis	None	None	No terms for herbicide or insect resistance	"GOSSYPIMUM BARBADENSE" "GOSSYPIMUM HIRSUTUM"	None
CABA	None	None	"HERBICIDE RESISTANCE" No terms for insect resistance	"GOSSYPIMUM BARBADENSE"/CT "GOSSYPIMUM HIRSUTUM"/CT	"TRANSGENIC PLANTS"

Database	Event	New proteins	Intended traits	Plant species	GM plants
CAS	None	None	"HERBICIDE RESISTANCE" No terms for insect resistance	GOSSYPIUM/CT	"GENETICALLY MODIFIED PLANTS"
Medline	None	None	"HERBICIDE RESISTANCE" No terms for insect resistance	"GOSSYPIUM BARBADENSE" "GOSSYPIUM HIRSUTUM"	"PLANTS, GENETICALLY MODIFIED"

The search results were limited to documents updated between July 1, 2022 and June 30, 2023 (UP>=20220701 and UP<=20230630), and to non-patent documents (not P/DT). To ensure that documents with indexing errors where two document types (DTs) (one eligible and one ineligible) were attached to a single record were not missed, documents with both 'journal' and 'patent' as document type were also kept. These putative documents would be identified with (P/DT AND J/DT) in CABA and CAPlus.

Table 4 summarizes the number of results obtained from each of the databases searched.

See [Appendix 2](#) for a complete search history.

Table 4: Overview of the selected databases and summary of search results from each database

Database	AGRICOLA	BIOSIS	CABA	CA-Plus	Medline
Database Provider	STN International	STN International	STN International	STN International	STN International
Coverage	1970-present	1926-present	1973-present	1907-present	1946-present
Date of search	14 Jul 2023	14 Jul 2023	14 Jul 2023	14 Jul 2023	14 Jul 2023
Datespan of the search	1 Jul 2022 – 30 Jun 2023	1 Jul 2022 – 30 Jun 2023	1 Jul 2022 – 30 Jun 2023	1 Jul 2022 – 30 Jun 2023	1 Jul 2022 – 30 Jun 2023
Latest database update	10 Jul 2023	12 Jul 2023	27 Jun 2023	13 Jul 2023	13 Jul 2023
Number of records retrieved	4	15	23	11	5
Number of records after duplicate removal	4	14	19	7	5

Database	AGRICOLA	BIOSIS	CABA	CA-Plus	Medline
Number of relevant records after rapid assessment	0	1	2	0	2

4. INTERNET and MANUAL SEARCHES

4.1. Internet Searches of food safety, agriculture, and biotechnology-related authority webpages

A search of the web pages of food safety, agriculture, and biotechnology-related authorities was conducted. Search results were manually examined for relevant records that were either published during the time period under consideration (date span of search: July 1, 2022 to June 30, 2023) or refer to relevant records published during this time frame. Relevance of results were determined based on the criteria listed in [Table 1](#) and they were summarized in Table 5. All web pages searched were justified by their recommendation in the EFSA 2019 explanatory note (1). Of the 13 key organisations cited in the EFSA 2019 explanatory note (1), Environment and Climate Change Canada and Intersecretarial Commission on Biosafety of GMOs (CIBIOGEM) were excluded, since they are not involved in the risk assessment of GM plants. The USDA, EPA, FDA, CFIA, Health Canada, FSANZ and MAFF websites were excluded, since these agencies do not regulate GM stacked products obtained by conventional breeding techniques. Therefore, the internet search was limited to four key organisations relevant for GHB614 x LLCotton25 x MON 15985 cotton. Search terms consisted of GHB614 x LLCotton25 x MON 15985 or BCS-GHØØ2-5 x ACS-GHØØ1-3 x MON-15985-7, 2mEPSPS or double mutant 5-enolpyruvyl shikimate-3-phosphate synthase enzyme, PAT/*bar* or Phosphinothricin, Cry1Ac, and Cry2Ab2 in GHB614 x LLCotton25 x MON 15985 cotton (all searched singly, with no search limits applied).

Table 5: Results of search of food safety, agriculture, and biotechnology-related authority websites

Source Site Name	Website URL	Date of Most Recent Site Update (dd.mm.yyyy)	Date of Search (dd.mm.yyyy)	No. of Relevant Records
Office of the Gene Technology Regulator (OGTR) Australia	http://www.ogtr.gov.au/	06.04.2023	10.07.2023	0
National Technical Commission on Biosafety (CTNBio) Brazil	http://ctnbio.mcti.gov.br/en	24.07.2023	17.-24.7.2023	0
National Advisory Commission on Agricultural Biotechnology (CONABIA) Argentina	https://www.argentina.gob.ar/agroindustria/bioeconomia/biotechnologia	17.07.2023	17.07.2023	0

Source Site Name	Website URL	Date of Most Recent Site Update (dd.mm.yyyy)	Date of Search (dd.mm.yyyy)	No. of Relevant Records
Genetic Engineering Approval Committee (GEAC) India	http://moef.gov.in/	19.07.2023	19.07.2023	0

4.2. Manual searches of reference lists of recent review articles

Recent review articles as sources of reference lists to search for potentially relevant studies were identified via searches of PubMed.gov for general terms such as “GMO” or “GM crops” in the titles and abstracts. The search of PubMed.gov was also restricted to recent reviews published between July 1, 2022 and June 30, 2023. The resulting number of relevant studies found within the bibliographies of these review articles is given in Table 6.

Table 6: Documents for which reference lists were scanned for relevant studies

No	Author(s) and Year	Title	Source	Number of relevant bibliographic references retrieved
1	Bhattacharjee S, Bhowmick R, Kant L, Paul K. 2023	Strategic transgene-free approaches of CRISPR-based genome editing in plants.	Mol Genet Genomics. 2023 May;298(3):507-520	0
2	Cermakova E, Lencova S, Mukherjee S, Horka P, Vobruba S, Demnerova K, Zdenkova K. 2023	Identification of Fish Species and Targeted Genetic Modifications Based on DNA Analysis: State of the Art.	Foods. 2023 Jan 3;12(1):228.	0
3	Connolly JB, Romeis J, Devos Y, Glandorf DCM, Turner G, Coulibaly MB. 2023	Gene drive in species complexes: defining target organisms.	Trends Biotechnol. 2023 Feb;41(2):154-164.	0
4	Eckerstorfer MF, Dolezel M, Engelhard M, Giovannelli V, Grabowski M, Heissenberger A, Lener M, Reichenbecher	Recommendations for the Assessment of Potential Environmental Effects of Genome-Editing Applications in Plants in the EU.	Plants (Basel). 2023 Apr 25;12(9):1764.	0

No	Author(s) and Year	Title	Source	Number of relevant bibliographic references retrieved
	W, Simon S, Staiano G, Wüst Saucy AG, Zünd J, Lüthi C. 2023			
5	Ghidoli M, Ponzoni E, Araniti F, Miglio D, Pilu R. 2023	Genetic Improvement of <i>Camelina sativa</i> (L.) Crantz: Opportunities and Challenges.	Plants (Basel). 2023 Jan 27;12(3):570.	0
6	Krasnodębski C, Sawuła A, Kaźmierczak U, Żuk M. 2023	Oligo-Not Only for Silencing: Overlooked Potential for Multidirectional Action in Plants.	Int J Mol Sci. 2023 Feb 24;24(5):4466.	0
7	Křížková B, Viktorová J, Lipov J. 2022	Approved Genetically Modified Potatoes (<i>Solanum tuberosum</i>) for Improved Stress Resistance and Food Safety.	J Agric Food Chem. 2022 Sep 28;70(38):11833-11843.	0
8	Liang J, Yang X, Jiao Y, Wang D, Zhao Q, Sun Y, Li Y, Wu K. 2022	The evolution of China's regulation of agricultural biotechnology.	aBIOTECH. 2022 Dec 5;3(4):237-249.	0
9	Platani M, Sokefun O, Bassil E, Apidianakis Y. 2023	Genetic engineering and genome editing in plants, animals, and humans: Facts and myths. Gene.	2023 Mar 10; 856:147141.	0
10	Pott A, Bundschuh M, Otto M, Schulz R. 2023	Assessing Effects of Genetically Modified Plant Material on the Aquatic Environment Using higher-tier Studies.	Bull Environ Contam Toxicol. 2023 Jan 2;110(1):35.	0
11	Rai GK, Kumar P, Choudhary SM, Kosser R, Khanday DM, Choudhary S, Kumar B, Magotra I, Kumar RR, Ram C, Roupheal Y, Corrado G, Behera TK. 2022	Biomimetic Strategies for Developing Abiotic Stress-Tolerant Tomato Cultivars:	An Overview. Plants (Basel). 2022 Dec 23;12(1):86	0

No	Author(s) and Year	Title	Source	Number of relevant bibliographic references retrieved
12	Rozas P, Kessi-Pérez EI, Martínez C. 2022	Genetically modified organisms: adapting regulatory frameworks for evolving genome editing technologies.	Biol Res. 2022 Oct 20;55(1):31.	0
13	Spök A, Sprink T, Allan AC, Yamaguchi T, Dayé C. 2022	Towards social acceptability of genome-edited plants in industrialised countries? Emerging evidence from Europe, United States, Canada, Australia, New Zealand, and Japan.	Front Genome Ed. 2022 Aug 31;4:899331.	0
14	Tatineni S, Hein GL. 2023	Plant Viruses of Agricultural Importance: Current and Future Perspectives of Virus Disease Management Strategies.	Phytopathology. 2023 Feb;113(2):117-141.	0
15	Tripathi S, Purchase D, Chandra R, Nadda AK, Bhargava PC. 2022	Mitigation of hazards and risks of emerging pollutants through innovative treatment techniques of post methanated distillery effluent - A review.	Chemosphere. 2022 Aug;300:134586 . doi: 10.1016/j.chemosphere.2022.134586.	0
16	Wang M, Wang H, Li K, Li X, Wang X, Wang Z. 2023	Review of CRISPR/Cas Systems on Detection of Nucleotide Sequences.	Foods. 2023 Jan 19;12(3):477.	0
17	Zimny T. 2023	Regulation of GMO field trials in the EU and new genomic techniques: will the planned reform facilitate experimenting with gene-edited plants?	BioTechnologia (Pozn). 2023 Mar 27;104(1):75-83.	0

5. RESULTS OF THE STUDY IDENTIFICATION AND SELECTION PROCESS

The database searches ([Section 0](#)) identified a total of 58 references, which were reduced to 49 references after removal of duplicates ([Table 4](#)). No additional studies were identified in the manual searches ([Section 4](#)).

5.1. Screening of titles and abstracts to exclude obviously irrelevant references (Stage 1)

All references identified in the database searches described in [Section 0](#) were assessed for relevance based on information in their title and abstract by two reviewers independently. If opinions of relevance differed, the discrepancies were discussed between the reviewers and if a disagreement persisted, the publication under discussion was transferred to Stage 2 for detailed evaluation by the experts. In this search, both evaluators were in 100% agreement.

Clearly irrelevant records were tagged as “Not Relevant”. These included:

- Duplicated entries
- Secondary literature (reviews), other than assessments from regulatory agencies
- Articles on non-relevant topics like detection methods, socio-economic implications of GM crops, GM policy, agronomical performance, other herbicide tolerant GM crops, other insect resistant GM crops, unrelated topics, etc.

Publications which appeared to be relevant and those of unclear relevance were tagged as “Relevant” and progressed to Stage 2 (detailed assessment; see [Section 5.2](#)).

The number of publications excluded after rapid assessment for relevance is presented in [Table 7](#) documenting the selection process.

5.2. Detailed assessment of eligible references (Stage 2)

Publications tagged as “Relevant” in Stage 1 were assessed in detail independently by two scientific experts in each of three corresponding areas (*i.e.*, Molecular Biology, Food and Feed Safety, Environmental Safety), based on the full text of the publications. If opinions of relevance differed between reviewers within each area, the initial reviewers discussed the discrepancy as necessary and consulted additional reviewers to resolve the discrepancy if needed.

Table 7 gives an overview of the reference selection process and results of the detailed assessment.

Table 7: Results of the publication selection process

Total number of publications retrieved after all searches of the scientific literature (excluding duplicates)	49
Number of publications excluded from the search results after rapid assessment for relevance (Stage 1)	44
Total number of full-text documents assessed in detail	5
Number of publications excluded from further consideration after detailed assessment for relevance (Stage 2)	5
Total number of unobtainable/unclear publications	0
Total number of relevant publications	0

Table 8 lists the publications determined to be relevant based on the detailed evaluation. Publications that were clearly not relevant after a detailed assessment are listed in Table 9. Table 10 lists the publications for which full-text documents were unobtainable for detailed assessment or for which relevance was unclear after detailed assessment.

Table 8: Report of all relevant publications retrieved after detailed assessment of full-text documents for relevance: ordered by category of information/data requirement(s)

Main category of information/data requirement	Study (Author(s) and year)	Title	Source
No publications in any category			

Table 9: Report of publications excluded from the risk assessment after detailed assessment of full-text documents

Study (Author(s) and year)	Title	Source	Reason(s) for exclusion based on eligibility/inclusion criteria listed in Table 1
Suassuna, N. D.; Morello, C. de L.; Perina, F. J.; Silva Filho, J. L. da; Pedrosa, M. B.; Magalhaes, F. O. da C.; Sofiat, V.; Lamas, F. M.; Chitarra, L. G.; Farias, F. J. C.; O, W. C. R. do; da Silva Filho, J. L.; do O, W. C. R. 2021	BRS 500 B2RF: transgenic cotton cultivar expressing Cry1Ac, Cry2Ab, and CP4-EPSPS with multiple disease resistance.	Crop Breeding and Applied Biotechnology (2021), Volume 21, Number 2, 12 refs. ISSN: 1518-7853; 1984-7033 DOI: https://doi.org/10.1590/1984-70332021v21n2c36 Published by: Brazilian Society of Plant Breeding, Londrina	This publication is a cultivar release related publication for the cultivar BRS 500 B2RF. GHB614 x LLCotton25 x MON 15985 cotton was not considered in this publication.

Study (Author(s) and year)	Title	Source	Reason(s) for exclusion based on eligibility/inclusion criteria listed in Table 1
Siddiqui Hamid Anees, Asad Shaheen, Naqvi Rubab Zahra, Asif Muhammad, Farooq Muhammad, Arshad Muhammad, Sarwar Muhammad, Amin Imran, Mukhtar Zahid, Mansoor Shahid, Liu Chengcheng Liu Xin, Abro Saifullah, Rizwan Muhammad 2022	Development and evaluation of triple gene transgenic cotton lines expressing three genes (Cry1Ac-Cry2Ab-EPSPS) for lepidopteran insect pests and herbicide tolerance .	Scientific reports, (20221101) Vol. 12, No. 1, pp. 18422. Electronic Publication Date: 1 Nov 2022 Journal code: 101563288. E-ISSN: 2045-2322. L-ISSN: 2045-2322. Report No.: PMC-PMC9626562.	The publication described development and evaluation of the molecular stack cotton plants NIBGE containing 3 gene cassettes for Cry1Ac-Cry2Ab-EPSPS. GHB614 x LLCotton25 x MON 15985 cotton was not considered in this publication.
Naegeli, H.; Bresson, J. L. Dalmay, T.; Dewhurst, I. C. Epstein, M. M.; Firbank, L. G. Guerche, P.; Hejatko, J. Moreno, F. J.; Mullins, E. Nogue, F.; Rostoks, N. Serrano, J. J. S.; Savoini, G. Veromann, E.; Veronesi, F. Alvarez, F.; Ardizzone, M. Raffaello, T. 2021	Assessment of genetically modified cotton GHB614 for renewal authorisation under regulation (EC) No 1829/2003 (application EFSA-GMO -RX-018).	EFSA Journal (2021), Volume 19, Number 7, 10 refs. ISSN: 1831-4732 DOI: https://doi.org/10.2903/j.efsa.2021.6671 Published by: Wiley, Oxford	This publication is about the EFSA evaluation of GHB614 cotton. Single events are not considered relevant for stacked events.

Study (Author(s) and year)	Title	Source	Reason(s) for exclusion based on eligibility/inclusion criteria listed in Table 1
Seixas Renato Nunes de Lima da Silveira Jose Maria Ferreira Jardim Ferrari Vinicius Eduardo 2022	Assessing environmental impact of genetically modified seeds in Brazilian agriculture.	Frontiers in bioengineering and biotechnology, (2022) Vol. 10, pp. 977793. Electronic Publication Date: 30 Aug 2022 Journal code: 101632513. ISSN: 2296-4185. L-ISSN: 2296-4185. Report No.: PMC-PMC9468974.	The authors used farm-level dataset originated from a survey conducted by a Céleres Consultancy, in Brazil. The survey collected data on production, revenue, costs, biotechnology adoption and pesticides used. Information on pesticide use was collected for harvest seasons 2009–2013 and covers 1,030 farms. No specific event was mentioned and no original data was presented in the publication. The ERA is not related to GHB614 x LLCotton25 x MON 15985 cotton.
Li Dongyang, Zhu Xiangzhen, Zhang Kaixin, Wang Li, Zhang Rui, Liang Chengzhen, Luo Junyu, Cui Jinjie 2022	Impact assessment of genetically modified herbicide -tolerant cotton on arthropod communities.	Journal of Cotton Research, (MAY 1 2022) Vol. 5, No. 1, pp. Article No.: 14. https://j cottonres.biomedcentral.com/ . ISSN: 2096-5044. E-ISSN: 2523-3254.	Arthropod communities were evaluated on the GM HT cotton variety GGK2 and its near-isogenic non-GM HT variety K312 in China. Arthropod composition was identical between GGK2 and K312. The ERA is not related to GHB614 x LLCotton25 x MON 15985 cotton.

Table 10: Report of unobtainable/unclear publications

Study (Author(s) and year)	Title	Source	Description of (unsuccessful) methods used to try and obtain a copy of the publication
No publications in this category.			

6. NARRATIVE SYNTHESIS/SUMMARY OF RELEVANT STUDIES

A total of five publications were selected during Stage 1 evaluation (rapid assessment based on title and abstract). After Stage 2 evaluation (detailed review based on full text), it was determined that none of the five publications were relevant for the safety assessment of the GHB614 x LLCotton25 x MON 15985 cotton and its newly expressed proteins, 2mEPSPS, PAT/*bar*, Cry1Ac, and Cry2Ab2.

7. CONCLUSION

The literature searches performed for the GHB614 x LLCotton25 x MON 15985 cotton and its newly expressed proteins, 2mEPSPS, PAT/*bar*, Cry1Ac, and Cry2Ab2, for the period from July 1, 2022 to June 30, 2023, identified a total of 49 unique publications (after duplicate removal). A total of five publication(s) were progressed for detailed assessment after excluding 44 obviously irrelevant publications during Stage 1 evaluation (rapid assessment based on title and abstract). The five publications that progressed to Stage 2 were evaluated in detail, based on full text, for potential relevance, following the pre-established criteria listed in [Table 1](#).

No new publications were found that contained new data on the molecular characterization of the GHB614 x LLCotton25 x MON 15985 cotton and its newly expressed proteins, 2mEPSPS, PAT/*bar*, Cry1Ac, and Cry2Ab2. Similarly, no new publications were found that suggested any potential adverse effects of GHB614 x LLCotton25 x MON 15985 cotton on human health, animal health, or the environment. No issues or topics were identified that would trigger or warrant more specific question formulation or indicate that a systematic review would be of value.

8. REFERENCES

No.	Author(s), title, source, edition, year, pages
1.	Devos Y, Guajardo IM, Alvarez F and Glanville J. Explanatory note on literature searching conducted in the context of GMO applications for (renewed) market authorisation and annual post-market environmental monitoring reports on GMOs authorised in the EU market. EFSA supporting publications 2019:EN-1614. 62 pages. doi:10.2903/sp.efsa.2019.EN-1614.

9. APPENDICES

Appendix 1: Database descriptions

Host	File	Description
STN	AGRICOLA	<p>Agriculture Online Access is a bibliographic database containing selected worldwide literature of agriculture and related fields. AGRICOLA is the locator and bibliographic access and control system of the National Agricultural Library (NAL) collections and also includes records from other cooperating institutions. Coverage of the database includes agricultural economics and rural sociology, agricultural production, animal sciences, chemistry, entomology, food and human nutrition, forestry, natural resources, pesticides, plant science, soils and fertilizers, and water resources. Also covered are related areas such as biology and biotechnology, botany, ecology, and natural history.</p> <p>The database draws on bibliographies, serial articles, book chapters, monographs, computer files, serials, maps, audiovisuals, and reports. Bibliographic information, abstracts, geographic terms, controlled terms, and supplementary terms are searchable.</p>
STN	BIOSIS	<p>BIOSIS Previews® is the largest and most comprehensive life science database in the world. Amongst others subject coverage includes Agriculture, Biochemistry, Biophysics, Botany, Environmental Biology, Physiology, Toxicology.</p> <p>Sources include periodicals, journals, conference proceedings, reviews, reports, patents, and short communications. Nearly 6,000 life source journals, 1,500 international meetings as well as review articles, books, and monographs are reviewed for inclusion.</p> <p>Bibliographic information, indexing terms, abstracts, and CAS Registry Numbers are all searchable.</p>
STN	CABA/CAB	<p>The CAB Abstracts database covers worldwide literature from all areas of agriculture and related sciences including Agriculture, Agricultural chemicals, Animal sciences and production, Crop protection, Crop sciences and production, Environment, Soils and fertilizers.</p> <p>Sources for CABA include journals, books, reports, published theses, conference proceedings, and patents.</p> <p>Bibliographic information, indexing terms, abstracts, and CAS Registry Numbers are searchable.</p>
STN	CAS/CAPLUS	<p>The Chemical Abstracts (CA) database covers all areas of Biochemistry, Chemistry and Chemical engineering, and related sciences.</p> <p>Sources include over 8,000 journals, patents from 38 national patent offices and two international patent organizations, technical reports, books, conference proceedings, and dissertations. Electronic only journals and Web preprints are also covered.</p> <p>Bibliographic terms, indexing terms, roles, CAS Registry Numbers, International Patent Classification, and abstracts are searchable.</p>

Host	File	Description
STN	MEDLINE	<p>MEDLINE contains information on every area of medicine. The MEDLINE database corresponds to Index Medicus, Index to Dental Literature, and International Nursing Index; OLDMEDLINE, with data from NLM's from the Cumulated Index Medicus (1960-1965) and Current List of Medical Literature (1958-1959); and, since August 2001, IN-PROCESS records, the latest documents before they have been completely indexed for inclusion on MEDLINE.</p> <p>Sources include journals and chapters in books or symposia. Bibliographic information, indexing terms, abstracts, chemical names, and CAS Registry Numbers are all searchable.</p> <p>Online thesauri are available for the Medical Subject Headings (/MN), Controlled Terms (/CT) and Chemical Name (/CN) fields.</p>

Appendix 2: Search history

```
FILE 'MEDLINE' ENTERED AT 08:11:08 ON 14 JUL 2023

L1      4 SEA GHB614 OR GHB(W)614 OR BCS-GH002-5 OR BCSGH002-5 OR
        BCS(W)GH002(W)5 OR BCSGH002(W)5 OR BCS-GH002-5 OR BCS(W)GH002(W)
        )5 OR BCSGH002(W)5

L2      0 SEA GHB614X OR GHB(W)614X OR BCS-GH002-5X OR BCSGH002-5X OR
        BCS(W)GH002(W)5X OR BCSGH002(W)5X OR BCS-GH002-5X OR BCS(W)GH00
        2(W)5X OR BCSGH002(W)5X

L3      4 SEA (L1 OR L2)

L4      3 SEA LLCOTTON25 OR LLCOTTON(W)25 OR LL(W)COTTON25 OR LL(W)COTTON
        (W)25 OR ACS-GH001-3 OR ACS(W)GH001(W)3 OR ACSGH001(W)3 OR
        ACS-GH001-3 OR ACS(W)GH001(W)3 OR ACSGH001(W)3

L5      0 SEA LLCOTTON25X OR LLCOTTON(W)25X OR LL(W)COTTON25X OR
        LL(W)COTTON(W)25X OR ACS-GH001-3X OR ACS(W)GH001(W)3X OR
        ACSGH001(W)3X OR ACS-GH001-3X OR ACS(W)GH001(W)3X OR ACSGH001(W)
        )3X

L6      0 SEA XLLCOTTON25 OR XLLCOTTON(W)25 OR XLL(W)COTTON25 OR
        XLL(W)COTTON(W)25 OR XACS-GH001-3 OR XACS(W)GH001(W)3 OR
        XACSGH001(W)3 OR XACS-GH001-3 OR XACS(W)GH001(W)3 OR XACSGH001(W)
        )3

L7      0 SEA XLLCOTTON25X OR XLLCOTTON(W)25X OR XLL(W)COTTON25X OR
        XLL(W)COTTON(W)25X OR XACS-GH001-3X OR XACS(W)GH001(W)3X OR
        XACSGH001(W)3X OR XACS-GH001-3X OR XACS(W)GH001(W)3X OR
        XACSGH001(W)3X

L8      3 SEA (L4 OR L5 OR L6 OR L7)

L9      12 SEA MON(W)15985 OR MON15985 OR MON-15985-7 OR MON(W)15985(W)7
        OR MON15985(W)7 OR XMON(W)15985 OR XMON15985 OR XMON-15985-7
        OR XMON(W)15985(W)7 OR XMON15985(W)7

L10     0 SEA GHB614XLLCOTTON25XMON15985 OR GHB(W)614XLLCOTTON25XMON15985
        OR GHB614XLL(W)COTTON25XMON15985 OR GHB614XLLCOTTON(W)25XMON15
        985 OR GHB614XLLCOTTON25XMON(W)15985 OR GHB(W)614XLL(W)COTTON25
        XMON15985 OR GHB(W)614XLLCOTTON(W)25XMON15985

L11     1 SEA GHB(W)614XLLCOTTON25XMON(W)15985 OR GHB(W)614XLL(W)COTTON(W)
        )25XMON15985 OR GHB(W)614XLL(W)COTTON25XMON(W)15986 OR
        GHB(W)614XLL(W)COTTON(W)25XMON(W)15985 OR GHB614(W)TIME#(W)LLCO
        TTON25(W)TIME#(W)MON15985 OR GHB614(W)TIME#(W)LLCOTTON25(W)TIME
        #(W)MON(W)15985

L12     1 SEA (L10 OR L11)

L13     1 SEA (L3 AND L8) OR (L3 AND L9) OR (L8 AND L9) OR L12

L14     2 SEA GLYTOL? OR GLYTOLTM? OR GLYTOLRTM? OR GLY(W)TOL? OR
        GLY(W)TOLTM? OR GLY(W)TOLRTM?

L15     73 SEA BOLLGARD? OR BOLGARD?

L16     18 SEA LIBERTYLINK OR LIBERTY(W)LINK OR LIBERTYLINKTM OR LIBERTY(W)
        )LINKTM OR LIBERTYLINKRTM OR LIBERTY(W)LINKRTM

L17     0 SEA (L14 AND L15) OR (L14 AND L16) OR (L15 AND L16)

L18     14 SEA 2MEPSPS OR 2(W)MEPSPS OR 2M(W)EPSPS OR 2(W)M(W)EPSPS

L19     4349 SEA EPSPS OR EPSP(W)SYNTHASE OR (ENOL(W)PYRUVYLSHIKIMATE OR
        ENOL(W)PYRUVYL(W)SHIKIMATE OR ENOLPYRUVYLSHIKIMATE OR ENOLPYRUV
        OYLSHIKIMATE OR ENOYLPRUVYOYLSHIKIMATE OR ENOLPYRUVYLSHIKIMIC) (
        4W) (PHOSPHATE OR PHOSPHORIC) (2W) (SYNTHASE OR SYNTHETASE)

L20     0 SEA (ENOLPYRUVYL OR ENOLPYRUYL OR ENOLPYRUVOYL) (W) (PHOSPHOSHIKI
        MATE OR PHOSPHOSHIKIMIC OR ENOLPYRUVYLSHIKIMATEPHOSPHATE) (2W) (S
        YNTHASE OR SYNTHETASE)

L21     421 SEA (ENOL(W)PYRUVYOYLSHIKIMATE OR ENOLPYRUVYLSHIKIMATE OR
        ENOLPYRUVYLSHIKIMIC OR ENOL(W) (PYRUVYL OR PYRUVOYL) (W) SHIKIMATE
        ) (3W) PHOSPHATE (W) (SYNTHASE OR SYNTHETASE)

L22     514 SEA (PHOSPHOSHIKIMATE (2W) CARBOXYVINYLTRANSFERASE OR PHOSPHOSHIK
        IMATE (2W) CARBOXYVINYL (W) TRANSFERASE OR ENOLPYRUVOYL (W) SHIKIMIC (
        3W) PHOSPHOSYNTHASE)

L23     25883 SEA ((DOUBL# OR DOBL#) (W) (MUTANT# OR MUTAT?) OR 2M)

L24     21 SEA L18 OR ((L19 OR L20 OR L21 OR L22)) (S) L23
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L25 1590 SEA ((BAR OR PAT) (2A) (GENE# OR PROTEIN# OR ENZYME#)) OR
PPT (2W) ACETYLTRANSFERASE OR PPT (2W) ACETYL (W) TRANSFERASE OR
PT (W) N (2W) ACETYLTRANSFERASE OR PT (W) N (2W) ACETYL (W) TRANSFERASE
L26 208 SEA PHOSPHINOTHRICIN (W) N (W) ACETYLTRANSFERASE OR PHOSPHINOTHRICI
N (2W) ACETYLTRANSFERASE OR PHOSPHINOTHRICIN (2W) ACETYL (W) TRANSFER
ASE OR PHOSPHINOTHRICINACETYL (W) TRANSFERASE
L27 1665 SEA (L25 OR L26)
L28 1284 SEA CRY1AC# OR CRY (W) 1 (W) AC# OR CRY (W) 1AC# OR CRY1 (W) AC# OR
CRY1 (W) A (W) C# OR CRY (W) 1 (W) A (W) C# OR CRY1A (W) C#
L29 127 SEA CRYIAC# OR CRY (W) I (W) AC# OR CRY (W) IAC# OR CRYI (W) AC# OR
CRYI (W) A (W) C# OR CRY (W) I (W) A (W) C# OR CRYIA (W) C#
L30 59 SEA CRYLAC# OR CRY (W) L (W) AC# OR CRY (W) LAC# OR CRYL (W) AC# OR
CRYL (W) A (W) C# OR CRY (W) L (W) A (W) C# OR CRYLA (W) C#
L31 94 SEA CRY2AB2 OR CRY (W) 2AB2 CRY2 (W) AB2 OR CRY2A (W) B2 OR CRY2AB (W)
2 OR CRY2 (W) AB (W) 2 OR CRY (W) 2 (W) AB2 OR CRY (W) 2 (W) A (W) B2 OR
CRYIIAB2 OR CRY (W) IIAB2 CRYII (W) AB2 OR CRYIIA (W) B2 OR CRYIIAB (W)
) 2 OR CRYII (W) AB (W) 2 OR CRY (W) II (W) AB2 OR CRY (W) II (W) A (W) B2
L32 1498 SEA (L28 OR L29 OR L30 OR L31)
L33 25 SEA (L24 AND L27) OR (L24 AND L32) OR (L27 AND L32)
L34 4025 SEA (HERBICID? OR GL!PHOSATE# OR GL!FOSATE# OR G360 OR G (W) 360
OR ROUNDUP? OR ROUND (W) UP?) (5A) (RESIST? OR TOLERAN? OR
PROTECT?)
L35 3818 SEA (HERBICID? OR BIALAPHOS OR BASTA OR GLUFOSINATE OR
GLUPHOSINATE OR PHOSPHINOTHRICIN OR LIBERTY?) (5A) (RESIST? OR
PROTECT? OR TOLERAN?)
L36 226792 SEA INSECT# OR PEST# OR LEPIDOPTERA# OR NOCTUIDAE OR CRAMBIDAE
OR BORER# OR CORNBORER# OR STALKBORER# OR EARWORM# OR EAR (W) WOR
M# OR ARMYWORM# OR ARMY (W) WORM# OR CUTWORM# OR CUT (W) WORM# OR
OSTRINIA OR O (W) NUBILALIS OR SESAMIA OR S (W) NONAGRIOIDES
L37 15162 SEA DIATRAEA OR D (W) GRANDIOSELLA OR D (W) CRAMBIDOIDES OR
HELICOVERPA OR H (W) ZEA OR SPODOPTERA OR S (W) FRUGIPERDA OR
PAPAIPEMA OR P (W) NEBRIS OR ELASMOPALPUS OR E (W) LIGNOSELLUS OR
D (W) SACCHARALIS OR STRIACOSTA OR S (W) ALBICOSTA
L38 26288 SEA AGROTIS OR A (W) IPSILON OR S (W) CRETICA OR MYTHIMNA OR
M (W) UNIPUNCTA OR ECB OR MCB OR SWCB OR SCSB OR CEW OR FAW OR
SCB OR WBC
L39 2693981 SEA (RESIST? OR PROTECT? OR TOLERAN?)
L40 9893 SEA ((L36 OR L37 OR L38)) (5A) L39
L41 3555 SEA (L34 AND L35) OR (L34 AND L40) OR (L35 AND L40)
L42 30477 SEA COTTON# OR GOSSYPIUM OR G (W) HIRSUTUM OR G (W) BARBADENSE
L43 4178777 SEA GMO OR GMOS OR LMO OR LMOS OR GM OR GE OR TRANSGEN? OR
(GENETIC? (3A) (MODIF? OR TRANSFORM? OR MANIPULAT? OR IMPROV? OR
ENGINEER?))
L44 121 SEA L41 AND L42 AND L43
L45 146 SEA L13 OR L17 OR L33 OR L44
L46 16 SEA L45 AND PY>=2021
L47 5 SEA L46 AND UP>=20220701 AND UP<=20230630

FILE 'BIOSIS' ENTERED AT 08:11:24 ON 14 JUL 2023

L48 5 SEA GHB614 OR GHB (W) 614 OR BCS-GH002-5 OR BCSGH002-5 OR
BCS (W) GH002 (W) 5 OR BCSGH002 (W) 5 OR BCS-GH002-5 OR BCS (W) GH002 (W)
) 5 OR BCSGH002 (W) 5
L49 0 SEA GHB614X OR GHB (W) 614X OR BCS-GH002-5X OR BCSGH002-5X OR
BCS (W) GH002 (W) 5X OR BCSGH002 (W) 5X OR BCS-GH002-5X OR BCS (W) GH00
2 (W) 5X OR BCSGH002 (W) 5X
L50 5 SEA (L48 OR L49)
L51 5 SEA LLCOTTON25 OR LLCOTTON (W) 25 OR LL (W) COTTON25 OR LL (W) COTTON
(W) 25 OR ACS-GH001-3 OR ACS (W) GH001 (W) 3 OR ACSGH001 (W) 3 OR
ACS-GH001-3 OR ACS (W) GH001 (W) 3 OR ACSGH001 (W) 3
L52 0 SEA LLCOTTON25X OR LLCOTTON (W) 25X OR LL (W) COTTON25X OR
LL (W) COTTON (W) 25X OR ACS-GH001-3X OR ACS (W) GH001 (W) 3X OR
ACSGH001 (W) 3X OR ACS-GH001-3X OR ACS (W) GH001 (W) 3X OR ACSGH001 (W)

) 3X
L53	0 SEA XLLCOTTON25 OR XLLCOTTON(W)25 OR XLL(W)COTTON25 OR XLL(W)COTTON(W)25 OR XACS-GH001-3 OR XACS(W)GH001(W)3 OR XACSGH001(W)3 OR XACS-GH001-3 OR XACS(W)GH001(W)3 OR XACSGH001(W)3
L54	0 SEA XLLCOTTON25X OR XLLCOTTON(W)25X OR XLL(W)COTTON25X OR XLL(W)COTTON(W)25X OR XACS-GH001-3X OR XACS(W)GH001(W)3X OR XACSGH001(W)3X OR XACS-GH001-3X OR XACS(W)GH001(W)3X OR XACSGH001(W)3X
L55	5 SEA (L51 OR L52 OR L53 OR L54)
L56	15 SEA MON(W)15985 OR MON15985 OR MON-15985-7 OR MON(W)15985(W)7 OR MON15985(W)7 OR XMON(W)15985 OR XMON15985 OR XMON-15985-7 OR XMON(W)15985(W)7 OR XMON15985(W)7
L57	0 SEA GHB614XLLCOTTON25XMON15985 OR GHB(W)614XLLCOTTON25XMON15985 OR GHB614XLL(W)COTTON25XMON15985 OR GHB614XLLCOTTON(W)25XMON15985 OR GHB614XLLCOTTON25XMON(W)15985 OR GHB(W)614XLL(W)COTTON25XMON15985 OR GHB(W)614XLLCOTTON(W)25XMON15985
L58	0 SEA GHB(W)614XLLCOTTON25XMON(W)15985 OR GHB(W)614XLL(W)COTTON(W)25XMON15985 OR GHB(W)614XLL(W)COTTON25XMON(W)15986 OR GHB(W)614XLL(W)COTTON(W)25XMON(W)15985 OR GHB614(W)TIME#(W)LLCOTTON25(W)TIME#(W)MON15985 OR GHB614(W)TIME#(W)LLCOTTON25(W)TIME#(W)MON(W)15985
L59	0 SEA (L57 OR L58)
L60	2 SEA (L50 AND L55) OR (L50 AND L56) OR (L55 AND L56) OR L59
L61	6 SEA GLYTOL? OR GLYTOLTM? OR GLYTOLRTM? OR GLY(W)TOL? OR GLY(W)TOLTM? OR GLY(W)TOLRTM?
L62	216 SEA BOLLGARD? OR BOLGARD?
L63	47 SEA LIBERTYLINK OR LIBERTY(W)LINK OR LIBERTYLINKTM OR LIBERTY(W)LINKTM OR LIBERTYLINKRTM OR LIBERTY(W)LINKRTM
L64	3 SEA (L61 AND L62) OR (L61 AND L63) OR (L62 AND L63)
L65	12 SEA 2MEPSPS OR 2(W)MEPSPS OR 2M(W)EPSPS OR 2(W)M(W)EPSPS
L66	5178 SEA EPSPS OR EPSP(W)SYNTHASE OR (ENOL(W)PYRUVYLSHIKIMATE OR ENOL(W)PYRUVYL(W)SHIKIMATE OR ENOLPYRUVYLSHIKIMATE OR ENOLPYRUVYLSHIKIMATE OR ENOYLPYRUVYLSHIKIMATE OR ENOLPYRUVYLSHIKIMIC) (4W) (PHOSPHATE OR PHOSPHORIC) (2W) (SYNTHASE OR SYNTHETASE)
L67	0 SEA (ENOLPYRUVYL OR ENOLPYRUYL OR ENOLPYRUVOYL) (W) (PHOSPHOSHIKIMATE OR PHOSPHOSHIKIMIC OR ENOLPYRUVYLSHIKIMATEPHOSPHATE) (2W) (SYNTHASE OR SYNTHETASE)
L68	761 SEA (ENOL(W)PYRUVYLSHIKIMATE OR ENOLPYRUVYLSHIKIMATE OR ENOLPYRUVYLSHIKIMIC OR ENOL(W) (PYRUVYL OR PYRUVOYL) (W)SHIKIMATE) (3W)PHOSPHATE(W) (SYNTHASE OR SYNTHETASE)
L69	29 SEA (PHOSPHOSHIKIMATE(2W)CARBOXYVINYLTRANSFERASE OR PHOSPHOSHIKIMATE(2W)CARBOXYVINYL(W)TRANSFERASE OR ENOLPYRUVOYL(W)SHIKIMIC(3W)PHOSPHOSYNTHASE)
L70	28113 SEA ((DOUBL# OR DOBL#) (W) (MUTANT# OR MUTAT?) OR 2M)
L71	21 SEA L65 OR (((L66 OR L67 OR L68 OR L69)) (S)L70)
L72	2987 SEA ((BAR OR PAT) (2A) (GENE# OR PROTEIN# OR ENZYME#)) OR PPT(2W)ACETYLTRANSFERASE OR PPT(2W)ACETYL(W)TRANSFERASE OR PT(W)N(2W)ACETYLTRANSFERASE OR PT(W)N(2W)ACETYL(W)TRANSFERASE
L73	337 SEA PHOSPHINOTHRICIN(W)N(W)ACETYLTRANSFERASE OR PHOSPHINOTHRICIN(2W)ACETYLTRANSFERASE OR PHOSPHINOTHRICIN(2W)ACETYL(W)TRANSFERASE OR PHOSPHINOTHRICINACETYL(W)TRANSFERASE
L74	3094 SEA (L72 OR L73)
L75	2058 SEA CRY1AC# OR CRY(W)1(W)AC# OR CRY(W)1AC# OR CRY1(W)AC# OR CRY1(W)A(W)C# OR CRY(W)1(W)A(W)C# OR CRY1A(W)C#
L76	231 SEA CRYIAC# OR CRY(W)I(W)AC# OR CRY(W)IAC# OR CRYI(W)AC# OR CRYI(W)A(W)C# OR CRY(W)I(W)A(W)C# OR CRYIA(W)C#
L77	367 SEA CRYLAC# OR CRY(W)L(W)AC# OR CRY(W)LAC# OR CRYL(W)AC# OR CRYL(W)A(W)C# OR CRY(W)L(W)A(W)C# OR CRYLA(W)C#
L78	131 SEA CRY2AB2 OR CRY(W)2AB2 CRY2(W)AB2 OR CRY2A(W)B2 OR CRY2AB(W)2 OR CRY2(W)AB(W)2 OR CRY(W)2(W)AB2 OR CRY(W)2(W)A(W)B2 OR CRYIIAB2 OR CRY(W)IIAB2 CRYII(W)AB2 OR CRYIIA(W)B2 OR CRYIIAB(W)

)2 OR CRYII(W)AB(W)2 OR CRY(W)II(W)AB2 OR CRY(W)II(W)A(W)B2
L79 2421 SEA (L75 OR L76 OR L77 OR L78)
L80 46 SEA (L71 AND L74) OR (L71 AND L79) OR (L74 AND L79)
L81 11863 SEA (HERBICID? OR GL!PHOSATE# OR GL!FOSATE# OR G360 OR G(W)360
OR ROUNDUP? OR ROUND(W)UP?) (5A) (RESIST? OR TOLERAN? OR
PROTECT?)
L82 10592 SEA (HERBICID? OR BIALAPHOS OR BASTA OR GLUFOSINATE OR
GLUPHOSINATE OR PHOSPHINOTHRICIN OR LIBERTY?) (5A) (RESIST? OR
PROTECT? OR TOLERAN?)
L83 1483914 SEA INSECT# OR PEST# OR LEPIDOPTERA# OR NOCTUIDAE OR CRAMBIDAE
OR BORER# OR CORNBORER# OR STALKBORER# OR EAWORM# OR EAR(W)WORM#
OR ARMYWORM# OR ARMY(W)WORM# OR CUTWORM# OR CUT(W)WORM# OR
OSTRINIA OR O(W)NUBILALIS OR SESAMIA OR S(W)NONAGRIOIDES
L84 29317 SEA DIATRAEA OR D(W)GRANDIOSELLA OR D(W)CRAMBIDOIDES OR
HELICOVERPA OR H(W)ZEA OR SPODOPTERA OR S(W)FRUGIPERDA OR
PAPAIPEMA OR P(W)NEBRIS OR ELASMOPALPUS OR E(W)LIGNOSELLUS OR
D(W)SACCHARALIS OR STRIACOSTA OR S(W)ALBICOSTA
L85 37155 SEA AGROTIS OR A(W)IPSILOIN OR S(W)CRETICA OR MYTHIMNA OR
M(W)UNIPUNCTA OR ECB OR MCB OR SWCB OR SCSB OR CEW OR FAW OR
SCB OR WBC
L86 2620674 SEA (RESIST? OR PROTECT? OR TOLERAN?)
L87 26819 SEA ((L83 OR L84 OR L85)) (5A)L86
L88 10225 SEA (L81 AND L82) OR (L81 AND L87) OR (L82 AND L87)
L89 79416 SEA COTTON# OR GOSSYPPIUM OR G(W)HIRSUTUM OR G(W)BARBADENSE
L90 495499 SEA GMO OR GMOS OR LMO OR LMOS OR GM OR GE OR TRANSGEN? OR
(GENETIC?(3A) (MODIF? OR TRANSFORM? OR MANIPULAT? OR IMPROV? OR
ENGINEER?))
L91 275 SEA L88 AND L89 AND L90
L92 326 SEA L60 OR L64 OR L80 OR L91
L93 29 SEA L92 AND PY>=2021
L94 15 SEA L93 AND UP>=20220701 AND UP<=20230630

FILE 'AGRICOLA' ENTERED AT 02:11:40 ON 14 JUL 2023
L95 2 SEA GHB614 OR GHB(W)614 OR BCS-GH002-5 OR BCSGH002-5 OR
BCS(W)GH002(W)5 OR BCSGH002(W)5 OR BCS-GH002-5 OR BCS(W)GH002(W)
5 OR BCSGH002(W)5
L96 0 SEA GHB614X OR GHB(W)614X OR BCS-GH002-5X OR BCSGH002-5X OR
BCS(W)GH002(W)5X OR BCSGH002(W)5X OR BCS-GH002-5X OR BCS(W)GH002(W)
5X OR BCSGH002(W)5X
L97 2 SEA (L95 OR L96)
L98 2 SEA LLCOTTON25 OR LLCOTTON(W)25 OR LL(W)COTTON25 OR LL(W)COTTON
(W)25 OR ACS-GH001-3 OR ACS(W)GH001(W)3 OR ACSGH001(W)3 OR
ACS-GH001-3 OR ACS(W)GH001(W)3 OR ACSGH001(W)3
L99 0 SEA LLCOTTON25X OR LLCOTTON(W)25X OR LL(W)COTTON25X OR
LL(W)COTTON(W)25X OR ACS-GH001-3X OR ACS(W)GH001(W)3X OR
ACSGH001(W)3X OR ACS-GH001-3X OR ACS(W)GH001(W)3X OR ACSGH001(W)
3X
L100 0 SEA XLLCOTTON25 OR XLLCOTTON(W)25 OR XLL(W)COTTON25 OR
XLL(W)COTTON(W)25 OR XACS-GH001-3 OR XACS(W)GH001(W)3 OR
XACSGH001(W)3 OR XACS-GH001-3 OR XACS(W)GH001(W)3 OR XACSGH001(W)
3
L101 0 SEA XLLCOTTON25X OR XLLCOTTON(W)25X OR XLL(W)COTTON25X OR
XLL(W)COTTON(W)25X OR XACS-GH001-3X OR XACS(W)GH001(W)3X OR
XACSGH001(W)3X OR XACS-GH001-3X OR XACS(W)GH001(W)3X OR
XACSGH001(W)3X
L102 2 SEA (L98 OR L99 OR L100 OR L101)
L103 9 SEA MON(W)15985 OR MON15985 OR MON-15985-7 OR MON(W)15985(W)7
OR MON15985(W)7 OR XMON(W)15985 OR XMON15985 OR XMON-15985-7
OR XMON(W)15985(W)7 OR XMON15985(W)7
L104 0 SEA GHB614XLLCOTTON25XMON15985 OR GHB(W)614XLLCOTTON25XMON15985
OR GHB614XLL(W)COTTON25XMON15985 OR GHB614XLLCOTTON(W)25XMON15985
OR GHB614XLLCOTTON25XMON(W)15985 OR GHB(W)614XLL(W)COTTON25

	XMON15985 OR GHB(W)614XLLCOTTON(W)25XMON15985
L105	0 SEA GHB(W)614XLLCOTTON25XMON(W)15985 OR GHB(W)614XLL(W)COTTON(W)25XMON15985 OR GHB(W)614XLL(W)COTTON25XMON(W)15986 OR GHB(W)614XLL(W)COTTON(W)25XMON(W)15985 OR GHB614(W)TIME#(W)LLCOTTON25(W)TIME#(W)MON15985 OR GHB614(W)TIME#(W)LLCOTTON25(W)TIME#(W)MON(W)15985
L106	0 SEA (L104 OR L105)
L107	0 SEA (L97 AND L102) OR (L97 AND L103) OR (L102 AND L103) OR L106
L108	3 SEA GLYTOL? OR GLYTOLTM? OR GLYTOLRTM? OR GLY(W)TOL? OR GLY(W)TOLTM? OR GLY(W)TOLRTM?
L109	203 SEA BOLLGARD? OR BOLGARD?
L110	42 SEA LIBERTYLINK OR LIBERTY(W)LINK OR LIBERTYLINKTM OR LIBERTY(W)LINKTM OR LIBERTYLINKRTM OR LIBERTY(W)LINKRTM
L111	3 SEA (L108 AND L109) OR (L108 AND L110) OR (L109 AND L110)
L112	4 SEA 2MEPSPS OR 2(W)MEPSPS OR 2M(W)EPSPS OR 2(W)M(W)EPSPS
L113	741 SEA EPSPS OR EPSP(W)SYNTHASE OR (ENOL(W)PYRUVYLSHIKIMATE OR ENOL(W)PYRUVYL(W)SHIKIMATE OR ENOLPYRUVYLSHIKIMATE OR ENOLPYRUVOYLSHIKIMATE OR ENOYLPYRUVOYLSHIKIMATE OR ENOLPYRUVYLSHIKIMIC) (4W) (PHOSPHATE OR PHOSPHORIC) (2W) (SYNTHASE OR SYNTHETASE)
L114	0 SEA (ENOLPYRUVYL OR ENOLPYRUYL OR ENOLPYRUVOYL) (W) (PHOSPHOSHIKIMATE OR PHOSPHOSHIKIMIC OR ENOLPYRUVYLSHIKIMATEPHOSPHATE) (2W) (SYNTHASE OR SYNTHETASE)
L115	337 SEA (ENOL(W)PYRUVOYLSHIKIMATE OR ENOLPYRUVYLSHIKIMATE OR ENOLPYRUVYLSHIKIMIC OR ENOL(W) (PYRUVYL OR PYRUVOYL) (W)SHIKIMATE) (3W)PHOSPHATE(W) (SYNTHASE OR SYNTHETASE)
L116	271 SEA (PHOSPHOSHIKIMATE(2W)CARBOXYVINYLTRANSFERASE OR PHOSPHOSHIKIMATE(2W)CARBOXYVINYL(W)TRANSFERASE OR ENOLPYRUVOYL(W)SHIKIMIC(3W)PHOSPHOSYNTHASE)
L117	7060 SEA ((DOUBL# OR DOBL#) (W) (MUTANT# OR MUTAT?) OR 2M)
L118	10 SEA L112 OR ((L113 OR L114 OR L115 OR L116)) (S)L117)
L119	829 SEA ((BAR OR PAT) (2A) (GENE# OR PROTEIN# OR ENZYME#)) OR PPT(2W)ACETYLTRANSFERASE OR PPT(2W)ACETYL(W)TRANSFERASE OR PT(W)N(2W)ACETYLTRANSFERASE OR PT(W)N(2W)ACETYL(W)TRANSFERASE
L120	258 SEA PHOSPHINOTHRICIN(W)N(W)ACETYLTRANSFERASE OR PHOSPHINOTHRICIN(2W)ACETYLTRANSFERASE OR PHOSPHINOTHRICIN(2W)ACETYL(W)TRANSFERASE OR PHOSPHINOTHRICINACETYL(W)TRANSFERASE
L121	911 SEA (L119 OR L120)
L122	1223 SEA CRY1AC# OR CRY(W)1(W)AC# OR CRY(W)1AC# OR CRY1(W)AC# OR CRY1(W)A(W)C# OR CRY(W)1(W)A(W)C# OR CRY1A(W)C#
L123	151 SEA CRYIAC# OR CRY(W)I(W)AC# OR CRY(W)IAC# OR CRYI(W)AC# OR CRYI(W)A(W)C# OR CRY(W)I(W)A(W)C# OR CRYIA(W)C#
L124	24 SEA CRYLAC# OR CRY(W)L(W)AC# OR CRY(W)LAC# OR CRYL(W)AC# OR CRYL(W)A(W)C# OR CRY(W)L(W)A(W)C# OR CRYLA(W)C#
L125	93 SEA CRY2AB2 OR CRY(W)2AB2 CRY2(W)AB2 OR CRY2A(W)B2 OR CRY2AB(W)2 OR CRY2(W)AB(W)2 OR CRY(W)2(W)AB2 OR CRY(W)2(W)A(W)B2 OR CRYIIAB2 OR CRY(W)IIAB2 CRYII(W)AB2 OR CRYIIA(W)B2 OR CRYIIAB(W)2 OR CRYII(W)AB(W)2 OR CRY(W)II(W)AB2 OR CRY(W)II(W)A(W)B2
L126	1441 SEA (L122 OR L123 OR L124 OR L125)
L127	21 SEA (L118 AND L121) OR (L118 AND L126) OR (L121 AND L126)
L128	9417 SEA (HERBICID? OR GL!PHOSATE# OR GL!FOSATE# OR G360 OR G(W)360 OR ROUNDUP? OR ROUND(W)UP?) (5A) (RESIST? OR TOLERAN? OR PROTECT?)
L129	8885 SEA (HERBICID? OR BIALAPHOS OR BASTA OR GLUFOSINATE OR GLUPHOSINATE OR PHOSPHINOTHRICIN OR LIBERTY?) (5A) (RESIST? OR PROTECT? OR TOLERAN?)
L130	355909 SEA INSECT# OR PEST# OR LEPIDOPTERA# OR NOCTUIDAE OR CRAMBIDAE OR BORER# OR CORNBORER# OR STALKBORER# OR EARWORM# OR EAR(W)WORM# OR ARMYWORM# OR ARMY(W)WORM# OR CUTWORM# OR CUT(W)WORM# OR OSTRINIA OR O(W)NUBILALIS OR SESAMIA OR S(W)NONAGRIOIDES
L131	15322 SEA DIATRAEA OR D(W)GRANDIOSELLA OR D(W)CRAMBIDOIDES OR HELICOVERPA OR H(W)ZEA OR SPODOPTERA OR S(W)FRUGIPERDA OR

PAPAIPEMA OR P(W)NEBRIS OR ELASMOPALPUS OR E(W)LIGNOSELLUS OR
D(W)SACCHARALIS OR STRIACOSTA OR S(W)ALBICOSTA
L132 8245 SEA AGROTIS OR A(W)IPSILON OR S(W)CRETICA OR MYTHIMNA OR
M(W)UNIPUNCTA OR ECB OR MCB OR SWCB OR SCSB OR CEW OR FAW OR
SCB OR WBC
L133 714744 SEA (RESIST? OR PROTECT? OR TOLERAN?)
L134 23346 SEA ((L130 OR L131 OR L132)) (5A)L133
L135 8722 SEA (L128 AND L129) OR (L128 AND L134) OR (L129 AND L134)
L136 68097 SEA COTTON# OR GOSSYPIMUM OR G(W)HIRSUTUM OR G(W)BARBADENSE
L137 113690 SEA GMO OR GMOS OR LMO OR LMOS OR GM OR GE OR TRANSGEN? OR
(GENETIC?(3A) (MODIF? OR TRANSFORM? OR MANIPULAT? OR IMPROV? OR
ENGINEER?))
L138 237 SEA L135 AND L136 AND L137
L139 259 SEA L107 OR L111 OR L127 OR L138
L140 11 SEA L139 AND PY>=2021
L141 4 SEA L140 AND UP>=20220701 AND UP<=20230630

FILE 'CABA' ENTERED AT 02:11:55 ON 14 JUL 2023
L142 7 SEA GHB614 OR GHB(W)614 OR BCS-GH002-5 OR BCSGH002-5 OR
BCS(W)GH002(W)5 OR BCSGH002(W)5 OR BCS-GH002-5 OR BCS(W)GH002(W)
5 OR BCSGH002(W)5
L143 0 SEA GHB614X OR GHB(W)614X OR BCS-GH002-5X OR BCSGH002-5X OR
BCS(W)GH002(W)5X OR BCSGH002(W)5X OR BCS-GH002-5X OR BCS(W)GH00
2(W)5X OR BCSGH002(W)5X
L144 7 SEA (L142 OR L143)
L145 5 SEA LLCOTTON25 OR LLCOTTON(W)25 OR LL(W)COTTON25 OR LL(W)COTTON
(W)25 OR ACS-GH001-3 OR ACS(W)GH001(W)3 OR ACSGH001(W)3 OR
ACS-GH001-3 OR ACS(W)GH001(W)3 OR ACSGH001(W)3
L146 0 SEA LLCOTTON25X OR LLCOTTON(W)25X OR LL(W)COTTON25X OR
LL(W)COTTON(W)25X OR ACS-GH001-3X OR ACS(W)GH001(W)3X OR
ACSGH001(W)3X OR ACS-GH001-3X OR ACS(W)GH001(W)3X OR ACSGH001(W)
3X
L147 0 SEA XLLCOTTON25 OR XLLCOTTON(W)25 OR XLL(W)COTTON25 OR
XLL(W)COTTON(W)25 OR XACS-GH001-3 OR XACS(W)GH001(W)3 OR
XACSGH001(W)3 OR XACS-GH001-3 OR XACS(W)GH001(W)3 OR XACSGH001(W)
3
L148 0 SEA XLLCOTTON25X OR XLLCOTTON(W)25X OR XLL(W)COTTON25X OR
XLL(W)COTTON(W)25X OR XACS-GH001-3X OR XACS(W)GH001(W)3X OR
XACSGH001(W)3X OR XACS-GH001-3X OR XACS(W)GH001(W)3X OR
XACSGH001(W)3X
L149 5 SEA (L145 OR L146 OR L147 OR L148)
L150 19 SEA MON(W)15985 OR MON15985 OR MON-15985-7 OR MON(W)15985(W)7
OR MON15985(W)7 OR XMON(W)15985 OR XMON15985 OR XMON-15985-7
OR XMON(W)15985(W)7 OR XMON15985(W)7
L151 0 SEA GHB614XLLCOTTON25XMON15985 OR GHB(W)614XLLCOTTON25XMON15985
OR GHB614XLL(W)COTTON25XMON15985 OR GHB614XLLCOTTON(W)25XMON15
985 OR GHB614XLLCOTTON25XMON(W)15985 OR GHB(W)614XLL(W)COTTON25
XMON15985 OR GHB(W)614XLLCOTTON(W)25XMON15985
L152 1 SEA GHB(W)614XLLCOTTON25XMON(W)15985 OR GHB(W)614XLL(W)COTTON(W)
)25XMON15985 OR GHB(W)614XLL(W)COTTON25XMON(W)15986 OR
GHB(W)614XLL(W)COTTON(W)25XMON(W)15985 OR GHB614(W)TIME#(W)LLCO
TTON25(W)TIME#(W)MON15985 OR GHB614(W)TIME#(W)LLCOTTON25(W)TIME
#(W)MON(W)15985
L153 1 SEA (L151 OR L152)
L154 2 SEA (L144 AND L149) OR (L144 AND L150) OR (L149 AND L150) OR
L153
L155 5 SEA GLYTOL? OR GLYTOLTM? OR GLYTOLRTM? OR GLY(W)TOL? OR
GLY(W)TOLTM? OR GLY(W)TOLRTM?
L156 402 SEA BOLLGARD? OR BOLGARD?
L157 105 SEA LIBERTYLINK OR LIBERTY(W)LINK OR LIBERTYLINKTM OR LIBERTY(W)
)LINKTM OR LIBERTYLINKRTM OR LIBERTY(W)LINKRTM
L158 6 SEA (L155 AND L156) OR (L155 AND L157) OR (L156 AND L157)

L159	14	SEA 2MEPSPS OR 2(W)MEPSPS OR 2M(W)EPSPS OR 2(W)M(W)EPSPS
L160	1268	SEA EPSPS OR EPSP(W)SYNTHASE OR (ENOL(W)PYRUVYLSHIKIMATE OR ENOL(W)PYRUVYL(W)SHIKIMATE OR ENOLPYRUVYLSHIKIMATE OR ENOLPYRUVYLSHIKIMATE OR ENOYL PYRUVYLSHIKIMATE OR ENOLPYRUVYLSHIKIMIC) (4W) (PHOSPHATE OR PHOSPHORIC) (2W) (SYNTHASE OR SYNTHETASE)
L161	0	SEA (ENOLPYRUVYL OR ENOLPYRUYL OR ENOLPYRUVOYL) (W) (PHOSPHOSHIKIMATE OR PHOSPHOSHIKIMIC OR ENOLPYRUVYLSHIKIMATEPHOSPHATE) (2W) (SYNTHASE OR SYNTHETASE)
L162	477	SEA (ENOL(W)PYRUVYLSHIKIMATE OR ENOLPYRUVYLSHIKIMATE OR ENOLPYRUVYLSHIKIMIC OR ENOL(W) (PYRUVYL OR PYRUVOYL) (W)SHIKIMATE) (3W) PHOSPHATE (W) (SYNTHASE OR SYNTHETASE)
L163	202	SEA (PHOSPHOSHIKIMATE (2W) CARBOXYVINYLTRANSFERASE OR PHOSPHOSHIKIMATE (2W) CARBOXYVINYL (W) TRANSFERASE OR ENOLPYRUVOYL (W) SHIKIMIC (3W) PHOSPHOSYNTHASE)
L164	7766	SEA ((DOUBL# OR DOBL#) (W) (MUTANT# OR MUTAT?) OR 2M)
L165	24	SEA L159 OR ((L160 OR L161 OR L162 OR L163)) (S) L164)
L166	1630	SEA ((BAR OR PAT) (2A) (GENE# OR PROTEIN# OR ENZYME#)) OR PPT (2W) ACETYLTRANSFERASE OR PPT (2W) ACETYL (W) TRANSFERASE OR PT (W) N (2W) ACETYLTRANSFERASE OR PT (W) N (2W) ACETYL (W) TRANSFERASE
L167	387	SEA PHOSPHINOTHRICIN (W) N (W) ACETYLTRANSFERASE OR PHOSPHINOTHRICIN (2W) ACETYLTRANSFERASE OR PHOSPHINOTHRICIN (2W) ACETYL (W) TRANSFERASE OR PHOSPHINOTHRICINACETYL (W) TRANSFERASE
L168	1736	SEA (L166 OR L167)
L169	2276	SEA CRY1AC# OR CRY (W) 1 (W) AC# OR CRY (W) 1AC# OR CRY1 (W) AC# OR CRY1 (W) A (W) C# OR CRY (W) 1 (W) A (W) C# OR CRY1A (W) C#
L170	236	SEA CRYIAC# OR CRY (W) I (W) AC# OR CRY (W) IAC# OR CRYI (W) AC# OR CRYI (W) A (W) C# OR CRY (W) I (W) A (W) C# OR CRYIA (W) C#
L171	43	SEA CRYLAC# OR CRY (W) L (W) AC# OR CRY (W) LAC# OR CRYL (W) AC# OR CRYL (W) A (W) C# OR CRY (W) L (W) A (W) C# OR CRYLA (W) C#
L172	158	SEA CRY2AB2 OR CRY (W) 2AB2 CRY2 (W) AB2 OR CRY2A (W) B2 OR CRY2AB (W) 2 OR CRY2 (W) AB (W) 2 OR CRY (W) 2 (W) AB2 OR CRY (W) 2 (W) A (W) B2 OR CRYIIAB2 OR CRY (W) IIAB2 CRYII (W) AB2 OR CRYIIA (W) B2 OR CRYIIAB (W) 2 OR CRYII (W) AB (W) 2 OR CRY (W) II (W) AB2 OR CRY (W) II (W) A (W) B2
L173	2631	SEA (L169 OR L170 OR L171 OR L172)
L174	38	SEA (L165 AND L168) OR (L165 AND L173) OR (L168 AND L173)
L175	20559	SEA (HERBICID? OR GL!PHOSATE# OR GL!FOSATE# OR G360 OR G(W) 360 OR ROUNDUP? OR ROUND (W) UP?) (5A) (RESIST? OR TOLERAN? OR PROTECT?)
L176	19754	SEA (HERBICID? OR BIALAPHOS OR BASTA OR GLUFOSINATE OR GLUPHOSINATE OR PHOSPHINOTHRICIN OR LIBERTY?) (5A) (RESIST? OR PROTECT? OR TOLERAN?)
L177	988145	SEA INSECT# OR PEST# OR LEPIDOPTERA# OR NOCTUIDAE OR CRAMBIDAE OR BORER# OR CORNBORER# OR STALKBORER# OR EARWORM# OR EAR (W) WORM# OR ARMYWORM# OR ARMY (W) WORM# OR CUTWORM# OR CUT (W) WORM# OR OSTRINIA OR O (W) NUBILALIS OR SESAMIA OR S (W) NONAGRIOIDES
L178	35287	SEA DIATRAEA OR D (W) GRANDIOSELLA OR D (W) CRAMBIDOIDES OR HELICOVERPA OR H (W) ZEA OR SPODOPTERA OR S (W) FRUGIPERDA OR PAPAIPEMA OR P (W) NEBRIS OR ELASMOPALPUS OR E (W) LIGNOSELLUS OR D (W) SACCHARALIS OR STRIACOSTA OR S (W) ALBICOSTA
L179	15101	SEA AGROTIS OR A (W) IPSILON OR S (W) CRETICA OR MYTHIMNA OR M (W) UNIPUNCTA OR ECB OR MCB OR SWCB OR SCSB OR CEW OR FAW OR SCB OR WBC
L180	1472770	SEA (RESIST? OR PROTECT? OR TOLERAN?)
L181	73942	SEA ((L177 OR L178 OR L179)) (5A) L180
L182	19420	SEA (L175 AND L176) OR (L175 AND L181) OR (L176 AND L181)
L183	112536	SEA COTTON# OR GOSSYPIMUM OR G (W) HIRSUTUM OR G (W) BARBADENSE
L184	204631	SEA GMO OR GMOS OR LMO OR LMOS OR GM OR GE OR TRANSGEN? OR (GENETIC? (3A) (MODIF? OR TRANSFORM? OR MANIPULAT? OR IMPROV? OR ENGINEER?))
L185	526	SEA L182 AND L183 AND L184
L186	564	SEA L154 OR L158 OR L174 OR L185
L187	35	SEA L186 AND PY>=2021

L188 23 SEA L187 AND UP>=20220701 AND UP<=20230630
L189 23 SEA L188 NOT P/DT
L190 0 SEA L188 AND (P/DT AND J/DT)
L191 23 SEA L189 OR L190

FILE 'HCAPLUS' ENTERED AT 02:12:19 ON 14 JUL 2023

L192 7 SEA GHB614 OR GHB(W)614 OR BCS-GH002-5 OR BCSGH002-5 OR
BCS(W)GH002(W)5 OR BCSGH002(W)5 OR BCS-GH002-5 OR BCS(W)GH002(W)
5 OR BCSGH002(W)5
L193 0 SEA GHB614X OR GHB(W)614X OR BCS-GH002-5X OR BCSGH002-5X OR
BCS(W)GH002(W)5X OR BCSGH002(W)5X OR BCS-GH002-5X OR BCS(W)GH002(W)5X
OR BCSGH002(W)5X
L194 7 SEA (L192 OR L193)
L195 11 SEA LLCOTTON25 OR LLCOTTON(W)25 OR LL(W)COTTON25 OR LL(W)COTTON
(W)25 OR ACS-GH001-3 OR ACS(W)GH001(W)3 OR ACSGH001(W)3 OR
ACSGH001-3 OR ACS(W)GH001(W)3 OR ACSGH001(W)3
L196 0 SEA LLCOTTON25X OR LLCOTTON(W)25X OR LL(W)COTTON25X OR
LL(W)COTTON(W)25X OR ACS-GH001-3X OR ACS(W)GH001(W)3X OR
ACSGH001(W)3X OR ACS-GH001-3X OR ACS(W)GH001(W)3X OR ACSGH001(W)
3X
L197 0 SEA XLLCOTTON25 OR XLLCOTTON(W)25 OR XLL(W)COTTON25 OR
XLL(W)COTTON(W)25 OR XACS-GH001-3 OR XACS(W)GH001(W)3 OR
XACSGH001(W)3 OR XACS-GH001-3 OR XACS(W)GH001(W)3 OR XACSGH001(W)
3
L198 0 SEA XLLCOTTON25X OR XLLCOTTON(W)25X OR XLL(W)COTTON25X OR
XLL(W)COTTON(W)25X OR XACS-GH001-3X OR XACS(W)GH001(W)3X OR
XACSGH001(W)3X OR XACS-GH001-3X OR XACS(W)GH001(W)3X OR
XACSGH001(W)3X
L199 11 SEA (L195 OR L196 OR L197 OR L198)
L200 33 SEA MON(W)15985 OR MON15985 OR MON-15985-7 OR MON(W)15985(W)7
OR MON15985(W)7 OR XMON(W)15985 OR XMON15985 OR XMON-15985-7
OR XMON(W)15985(W)7 OR XMON15985(W)7
L201 0 SEA GHB614XLLCOTTON25XMON15985 OR GHB(W)614XLLCOTTON25XMON15985
OR GHB614XLL(W)COTTON25XMON15985 OR GHB614XLLCOTTON(W)25XMON15
985 OR GHB614XLLCOTTON25XMON(W)15985 OR GHB(W)614XLL(W)COTTON25
XMON15985 OR GHB(W)614XLLCOTTON(W)25XMON15985
L202 1 SEA GHB(W)614XLLCOTTON25XMON(W)15985 OR GHB(W)614XLL(W)COTTON(W)
)25XMON15985 OR GHB(W)614XLL(W)COTTON25XMON(W)15986 OR
GHB(W)614XLL(W)COTTON(W)25XMON(W)15985 OR GHB614(W)TIME#(W)LLCO
TTON25(W)TIME#(W)MON15985 OR GHB614(W)TIME#(W)LLCOTTON25(W)TIME
#(W)MON(W)15985
L203 1 SEA (L201 OR L202)
L204 9 SEA (L194 AND L199) OR (L194 AND L200) OR (L199 AND L200) OR
L203
L205 8 SEA GLYTOL? OR GLYTOLTM? OR GLYTOLRTM? OR GLY(W)TOL? OR
GLY(W)TOLTM? OR GLY(W)TOLRTM?
L206 190 SEA BOLLGARD? OR BOLGARD?
L207 53 SEA LIBERTYLINK OR LIBERTY(W)LINK OR LIBERTYLINKTM OR LIBERTY(W)
)LINKTM OR LIBERTYLINKRTM OR LIBERTY(W)LINKRTM
L208 4 SEA (L205 AND L206) OR (L205 AND L207) OR (L206 AND L207)
L209 31 SEA 2MEPSPS OR 2(W)MEPSPS OR 2M(W)EPSPS OR 2(W)M(W)EPSPS
L210 4631 SEA EPSPS OR EPSP(W)SYNTHASE OR (ENOL(W)PYRUVYLSHIKIMATE OR
ENOL(W)PYRUVYL(W)SHIKIMATE OR ENOLPYRUVYLSHIKIMATE OR ENOLPYRUV
OYLSHIKAMATE OR ENOYLPRUVOYLSHIKIMATE OR ENOLPYRUVYLSHIKIMIC) (4W)
(PHOSPHATE OR PHOSPHORIC) (2W) (SYNTHASE OR SYNTHETASE)
L211 9 SEA (ENOLPYRUVYL OR ENOLPYRUYL OR ENOLPYRUVOYL) (W) (PHOSPHOSHIKI
MATE OR PHOSPHOSHIKIMIC OR ENOLPYRUVYLSHIKIMATEPHOSPHATE) (2W) (S
YNTHASE OR SYNTHETASE)
L212 1133 SEA (ENOL(W)PYRUVOYLSHIKIMATE OR ENOLPYRUVYLSHIKIMATE OR
ENOLPYRUVYLSHIKIMIC OR ENOL(W) (PYRUVYL OR PYRUVOYL) (W) SHIKIMATE
) (3W) PHOSPHATE(W) (SYNTHASE OR SYNTHETASE)
L213 90 SEA (PHOSPHOSHIKIMATE(2W)CARBOXYVINYLTRANSFERASE OR PHOSPHOSHIK

IMATE (2W) CARBOXYVINYL (W) TRANSFERASE OR ENOLPYRUVYL (W) SHIKIMIC (3W) PHOSPHOSYNTHASE)

L214 77236 SEA ((DOUBL# OR DOBL#) (W) (MUTANT# OR MUTAT?) OR 2M)

L215 46 SEA L209 OR ((L210 OR L211 OR L212 OR L213)) (S) L214)

L216 5706 SEA ((BAR OR PAT) (2A) (GENE# OR PROTEIN# OR ENZYME#)) OR PPT (2W) ACETYLTRANSFERASE OR PPT (2W) ACETYL (W) TRANSFERASE OR PT (W) N (2W) ACETYLTRANSFERASE OR PT (W) N (2W) ACETYL (W) TRANSFERASE

L217 809 SEA PHOSPHINOTHRICIN (W) N (W) ACETYLTRANSFERASE OR PHOSPHINOTHRICIN (2W) ACETYLTRANSFERASE OR PHOSPHINOTHRICIN (2W) ACETYL (W) TRANSFERASE OR PHOSPHINOTHRICINACETYL (W) TRANSFERASE

L218 6023 SEA (L216 OR L217)

L219 2451 SEA CRY1AC# OR CRY (W) 1 (W) AC# OR CRY (W) 1AC# OR CRY1 (W) AC# OR CRY1 (W) A (W) C# OR CRY (W) 1 (W) A (W) C# OR CRY1A (W) C#

L220 1990 SEA CRYIAC# OR CRY (W) I (W) AC# OR CRY (W) IAC# OR CRYI (W) AC# OR CRYI (W) A (W) C# OR CRY (W) I (W) A (W) C# OR CRYIA (W) C#

L221 74 SEA CRYLAC# OR CRY (W) L (W) AC# OR CRY (W) LAC# OR CRYL (W) AC# OR CRYL (W) A (W) C# OR CRY (W) L (W) A (W) C# OR CRYLA (W) C#

L222 168 SEA CRY2AB2 OR CRY (W) 2AB2 CRY2 (W) AB2 OR CRY2A (W) B2 OR CRY2AB (W) 2 OR CRY2 (W) AB (W) 2 OR CRY (W) 2 (W) AB2 OR CRY (W) 2 (W) A (W) B2 OR CRYIIAB2 OR CRY (W) IIAB2 CRYII (W) AB2 OR CRYIIA (W) B2 OR CRYIIAB (W) 2 OR CRYII (W) AB (W) 2 OR CRY (W) II (W) AB2 OR CRY (W) II (W) A (W) B2

L223 3280 SEA (L219 OR L220 OR L221 OR L222)

L224 3 SEA (L205 AND L208) OR (L205 AND L213) OR (L208 AND L213)

L225 32162 SEA (HERBICID? OR GL!PHOSATE# OR GL!FOSATE# OR G360 OR G (W) 360 OR ROUNDUP? OR ROUND (W) UP?) (5A) (RESIST? OR TOLERAN? OR PROTECT?)

L226 30987 SEA (HERBICID? OR BIALAPHOS OR BASTA OR GLUFOSINATE OR GLUPHOSINATE OR PHOSPHINOTHRICIN OR LIBERTY?) (5A) (RESIST? OR PROTECT? OR TOLERAN?)

L227 342997 SEA INSECT# OR PEST# OR LEPIDOPTERA# OR NOCTUIDAE OR CRAMBIDAE OR BORER# OR CORNBORER# OR STALKBORER# OR EARWORM# OR EAR (W) WORM# OR ARMYWORM# OR ARMY (W) WORM# OR CUTWORM# OR CUT (W) WORM# OR OSTRINIA OR O (W) NUBILALIS OR SESAMIA OR S (W) NONAGRIOIDES

L228 25584 SEA DIATRAEA OR D (W) GRANDIOSELLA OR D (W) CRAMBIDOIDES OR HELICOVERPA OR H (W) ZEA OR SPODOPTERA OR S (W) FRUGIPERDA OR PAPAIPEMA OR P (W) NEBRIS OR ELASMOPALPUS OR E (W) LIGNOSELLUS OR D (W) SACCHARALIS OR STRIACOSTA OR S (W) ALBICOSTA

L229 29594 SEA AGROTIS OR A (W) IPSILON OR S (W) CRETICA OR MYTHIMNA OR M (W) UNIPUNCTA OR ECB OR MCB OR SWCB OR SCSB OR CEW OR FAW OR SCB OR WBC

L230 6571734 SEA (RESIST? OR PROTECT? OR TOLERAN?)

L231 44791 SEA ((L227 OR L228 OR L229)) (5A) L230

L232 30525 SEA (L225 AND L226) OR (L225 AND L231) OR (L226 AND L231)

L233 293563 SEA COTTON# OR GOSSYPIUM OR G (W) HIRSUTUM OR G (W) BARBADENSE

L234 747660 SEA GMO OR GMOS OR LMO OR LMOS OR GM OR GE OR TRANSGEN? OR (GENETIC? (3A) (MODIF? OR TRANSFORM? OR MANIPULAT? OR IMPROV? OR ENGINEER?))

L235 1402 SEA L232 AND L233 AND L234

L236 1410 SEA L204 OR L208 OR L224 OR L235

L237 335 SEA L236 AND PY>=2021

L238 64 SEA L237 AND UP>=20220701 AND UP<=20230630

L239 11 SEA L238 NOT P/DT

L240 0 SEA L238 AND (P/DT AND J/DT)

L241 11 SEA L239 OR L240

FILE 'MEDLINE, BIOSIS, AGRICOLA, CABA, HCAPLUS' ENTERED AT 02:12:35 ON 14
JUL 2023
L242 49 DUP REM L47 L94 L141 L191 L241 (9 DUPLICATES REMOVED)
ANSWERS '1-5' FROM FILE MEDLINE
ANSWERS '6-19' FROM FILE BIOSIS
ANSWERS '20-23' FROM FILE AGRICOLA
ANSWERS '24-42' FROM FILE CABA
ANSWERS '43-49' FROM FILE HCAPLUS